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AX698788
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ACCESSION: AXO98788
ACCESSION: AXA770435
ACCESSION: AXA770746
ACCESSION: AX624408
ACCESSION: AX624704
ACCESSION: AX624745
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ACCESSION: AX624234
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ACCESSION: AX826542
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RESULT 2 AX599526/c LOCUS LOCUS DEFINITION ACCESSION VERSION VERSION KEYWORDS SOURCE ORGANISM REFERENCE AUTHORS TITLE JOURNAL FEATURES SOURCE	Best Local Matches 1  Qy 2079  Db 16	Truce 5% S	0 0 3 5 5 4 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9
AX599526 Sequence 866 from Pate, AX599526 AX599526.1 GI:2839967 Extificial sequences. Berlin,K., Braun,A., Dolek,A., Piepenbrock,C. Lewin,A., Lipscher,E., Peletods and nucleic ac proliferative disorder Patent: WO 0207727-A Epigenomics AG (DE) Epigenomics AG (DE) Corganiam="sy/mol_type="un/db_xref="texx/note="Detect"	Similarity 93.8 15; Conservative 79 TCCAAAGAAACACTAA	Q807902 Q807902. Q807902. Q807902. rtificia rtificia oekens,J immrich, arx,A. ethod an rolifera atent: W pigenomi	. 4 4 4 4 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
ioan i se	%; Pre 0; 2094	from Patent :47113296 truct truct uences. ujan,T., Ko ujan,T., Sch disorders disorders fd1803-A 11 (01803-A 11 (01803)-A 1	11 1 AX626388 11 1 AX627142 11 1 AX627434 11 1 AX627701 11 1 AX631829 11 1 AX632125 11 1 AX632125 11 1 BD124358 ALIGNI
18 bp DNA ent w002077272.  70  Distler, J., Guetig, D. C., Adorjan, P., Grabs, Maier, S., Model, F., H. cids for the analysis s6 03-OCT-2002; lifiers lifiers ynthetic construct" nassigned DNA" xon:32630"	. No. 2.2 ismatches	DNA 4035803 4035803 T., Mai A., Sch A., Sch ce impro -APR-20 -APR-20 DNA"	26388 267142 277434 27701 27701 32129 32125 32126 24358 ALIGNMENTS
linear PAT 14-FEB-2003  , Howe, A., Mueller, J., ,G., Lesche, R., Leu, E., , Mueller, V., Otto, T.,  of hematopoietic cell  for CDKN1C"	TO TO	linear	ACCESSION: AX626388 ACCESSION: AX627142 ACCESSION: AX627434 ACCESSION: AX627701 ACCESSION: AX631829 ACCESSION: AX632125 ACCESSION: AX632156 ACCESSION: BD124358

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Sequence 508 from Patent WO03044226.
AX767860
                                                                                                                                                                                                         Method and nucleic acids for the analysis of a lung cell proliferative disorder
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synthetic construct
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                                                                                                                                                                                              Patent: WO 03052135-A 781 26-JUN-2003;
                                                                                                                                                                                                                                     and Nimmrich, I.
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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                                                                                                    note="Detection oligonucleotide for CDKN1C"
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/mol_type="unassigned DNA"
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AX822902
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Sequence 794 from Patent
AX826542
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AR056349/c
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Sequence 553 from patent US 5
AR056349
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Sequence 3813 from patent
AR326411
                                                                                                                                                                                                                                                                                                                                                                                                                                           1 (bases 1 to 17) Pavco, P., McSwiggen, J.A., Stinchcomb, D.T. and Escobedo, J. Method and reagent for the treatment of diseases or conditions related to levels of vascular endothelial growth factor receptor Patent: US 6566127-A 3813 20-MAY-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 (bases 1 to 17)
Pavco, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6346398-A 8030 12-FEB-2002;
Location/Qualifiers
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                                                                              Draper, K.G
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Grimm,S., Stinchcomb,D.T., McSwiggen,J., Sullivan,S. and
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                                                   Intercellular adhesion molecule-1 (ICAM-1) ribozymes
Patent: US 5837542-A 553 17-NOV-1998;
                                                                                                                      Unclassified
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            organism="unknown"
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AR114107
                                                                                                                                                                                                                                 Stinchcomb,D.T., Dudyzz,L.W., Chowrira,B., Grimm,S., Direnzo,A., Karpeisky,A., Draper,K.G., Kisich,K., Matulic-Adamic,J., McKarpeisky,A., Modak,A., Pavco,P., Beigelman,L., Sullivan,S.M., Sweedler,D., Thompson,J.D., Tracz,D., Usman,N., Wincott,F.E. and
                                                                                                                                                                                                                                                                                                                                                                     Sequence 713 from Patent EP1260586. AX633574
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ribozyme treatment of diseases or conditions related to levels intercellular adhesion molecule-1 (ICAM-1) Patent: US 6132967-A 553 17-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 (Dases 1 to 15)
Grimm,S., Stinchcomb,D.T., McSwiggen,J., Sullivan,S.
Draper,K.G.
                                                                                                                                                    Patent: EP 1260586-A 713 27-NOV-2002; RIBOZYME PHARMACEUTICALS, INC. (US) Location/Qualifiers
                                                                                                                                                                                                                                                                                                                 unidentified
unidentified
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                                                                                                                                                                                                                        Woolf, T.
                                                                                                                                                                                                                                                                                                   unclassified
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553 from patent US 6132967.
                                                                                                /organism="unidentified"
/mol_type="unassigned RNA"
/db_xref="taxon:32644"
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/mol_type="unassigned DNA"
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Method for determining the homeostasis of hairy skin
Patent: WO 2004059002-A 561 15-JUL-2004;
Henkel Kommanditgesellschaft auf Aktien (DE)
Location/Qualifiers
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1
                                                       Homo sapiens (human)
                                                                                  AX624234.1
                                                                                              Sequence 1275 from Patent AX624234
                                                                                                                          AX624234
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Zoghbi, H.Y., Van den Veyver, I.B., Amir, R. and Francke, U.

Rethod of screening Rett syndrome by detecting a mutation in MECP2

Patent: US 6709817-A 94 23-MAR-2004;
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Sequence 561 from Patent WO2004059002.
CQ833190
CQ833190.1 GI:50832797
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                       /organism="unknown"
/mol_type="genomic DNA"
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Sequence 8697 from Patent
AX631655
                                                                                                       Petersohn,D., Conradt,M. and Hofmann,K. Method for determining homeostasis of the Patent: WO 2053774-A 8697 11-JUL-2002; Henkel Kommanditgesellschaft auf Aktien (D
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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CQ832709/c
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Wang,C.-G. and Hepburn,A.G.
Genetic sequence assay using DNA triple strand formation
Patent: US 5861244-A 274 19-JAN-1999;
                                                                                                                                                                                                                                                                    Conradt, M. and Hofmann, K.
Method for determining the homeostasis of hairy skin
Patent: WO 2004059002-A 80 15-JUL-2004;
Henkel Kommanditgesellschaft auf Aktien (DE)
Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Conservative 0; Mismatches
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Pred. No. 21;
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                                          Homo sapiens (human)
Homo sapiens
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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Sequence 1306 from Patent
CQ833935
CQ833935.1 GI:50833542
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Method for determining the homeostasis of hairy skin Patent: WO 2004059002-A 423 15-JUL-2004;
Henkel Kommanditgesellschaft auf Aktien (DE)
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Conradt,M. and Hofmann,K.
Method for determining the homeostasis of hairy skin
Patent: WO 2004059002-A 417 15-JUL-2004;
Henkel Kommanditgesellschaft auf Aktien (DE)
Location/Qualifiers
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Homo sapiens
Petersohn, D., Schlotmann, K., Conradt, M. and Hofmann, K.
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Homo sapiens
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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WO2004059002
                 Gassenmeier, T., Holtkoetter, O.,
                                          Craniata; Vertebrata; Catarrhini; Hominidae;
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                                                         Euteleostomi;
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                                                                                                                                                                                                             Sequence 189 from patent US 6
AR301608
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Method for determining markers of human facial skin
Patent: WO 2004059001-A 287 15-JUL-2004;
Henkel Kommanditgesellschaft auf Aktien (DE)
Location/Qualifiers
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                                                                                      Compositions and methods for wound healing Patent: US 6538173-A 189 25-MAR-2003;
Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Heber-Katz,E.
                                                                                                                                                                                                     AR301608.1 GI:31689410
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/mol_type="unassigned DN/
/db_xref="taxon:9606"
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/mol_type="unassigned DNJ/db_xref="taxon:9606"
                                               /organism="unknown"
/mol_type="genomic
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Dutta,S., Biswas,B. and Vemulapalli,R.
Size-variable strain-specific protective antigen for potomac horse
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AR367551
AR367551.1 GI:34600862
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Dutta, S., Biswas, B. and Vemulapalli, R.
Size-variable strain-specific protecti
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Location/Qualifiers
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/mol_type="genomic DNA"
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/mol_type="genomic |
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from patent US 6375954.
                                            GI:13538028
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Polymorphisms in the human cyp3a4 and cyp3a7 genes and their use
diagnostic and therapeutic applications
Patent: WO 0120025-A 94 22-MAR-2001;
Epidauros Biotechnologie AG (DE)
Location/Qualifiers
                                                                                                             Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 12 from Patent W002053773.
AX470435 AX470435.1 GI:22205560
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                                                        Hofmann, K., Conradt, M. and Petersohn, D. Method for determining skin stress or skin ageing Patent: WO 02053773-A 12 11-JUL-2002;
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/mol type="unassigned DNA"
/db_xref="texon.32630"
/note="artificial"
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
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organism="Homo sapiens"/
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AX624408/c
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AX470746
AX470746.1 GI:22205871
                                                                                                                              Method for determining homeostasis of the Patent: WO 02033774-A 1449 11-JUL-2002; Henkel Kommanditgesellschaft auf Aktien Location/Qualifiers
                                                                                                                                                                                                                                                                                      Sequence 1449 from Patent AX624408
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hofmann,K., Conradt,M. and Petersohn,D. Method for determining skin stress or skin Patent: WO 02053773-A 323 11-JUL-2002; HENKEL KGAA (DE)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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/db_xref="taxon:9606"
                                                                           /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Sequence 3419 from Patent
AX626378
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Sequence 1786 from Patent
AX624745
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              Homo sapiens (human)
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Patent: WO 02053774-A 1786 11-JUL-2002;
Henkel Kommanditgesellschaft auf Aktien (DE)
Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/db_xref="taxon:9606"
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                                                                                                                                                Homo sapiens (human)
Homo sapiens
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Sequence 4183 from Patent WO02053774.
AX627142 GI:28455180
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Sequence 3429 from Patent
AX626388
AX626388.1 GI:28454426
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                                                                         Petersohn,D., Conradt,M. and Hofmann,K. Method for determining homeostasis of the skin Patent: WO 02053794-A 4183 11-UUL-2002; Henkel Kommanditgesellschaft auf Aktien (DE)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNF
/db_xref="taxon:9606"
/organism="Homo sapiens"
/mol_type="unassigned DNA"
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Seguence 4475 from Patent
AX627434
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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RESULT 38

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AX632125/c
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AX632166
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AX632166
AX632166.1
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AX632125
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AX631829
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           Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Method for determining homeostasis of the Patent: WO 02053774-A 8871 11-JUL-2002; Henkel Kommanditgesellschaft auf Aktien
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Mammalia; Eutheria; Primates;
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/mol_type="unassigned DN/
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Best Local Similarity 90.9
Matches 10; Conservative
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Best Local Similarity 90.9%;
Matches 10; Conservative
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                                                                                   2087 AACACTAAGCT 2097
                                                                                                                                                                                                                                                                                                                                                                                                       Compositions and method for healing wound Patent: JP 2002503460-A 189 05-FEB-2002;
THE WISTAR INSTITUTE
OS Mus musculus (mouse)
PN JP 2002503460-A/189
PD JP-EB-199 JP 200531545
PF 12-FEB-199 US 60/074737,26-AUG-1998
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PR 13-FEB-1998 US 60/074737,26-AUG-1998 US 60/097937 PR
28-SEP-1998 US 60/102051
PI ELLEN HEBER KATZ
PC C12N15/09,A01K67/027,C12N5/10,C12Q1/68,G01N33/50,C12N15/00, PC
C12N5/00
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Mus musculus (house mouse)
Mus musculus
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                                                                   AACACCAAGCT 11
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Location/Qualifiers
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/organism='Mus musculus (mouse)'.
Location/Qualifiers
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Copyright (c) 1993 - 2004 Compugen Ltd.
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     DB
US-08-584-040-8030
US-09-977772B-3813
US-08-292-620A-553
US-09-071-845-53
US-09-071-845-53
US-09-38-353-415
US-08-388-551B-415
US-08-488-551B-702
US-08-488-551B-702
US-09-157-013-94
US-09-157-257-17
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US-09-18-383-33
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US-08-388-353-703
US-08-488-551B-414
US-08-488-551B-414
US-08-488-551B-414
US-08-488-551B-703
US-09-677-1288-50
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Sequence 8030, Ap
Sequence 3813, App
Sequence 553, App
Sequence 415, App
Sequence 702, App
Sequence 702, App
Sequence 94, Appl
Sequence 274, App
Sequence 17, Appl
Sequence 189, Appl
Sequence 189, Appl
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Sequence 32, Appl
Sequence 33, Appl
Sequence 31, Appl
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Sequence 701, App
Sequence 50, Appl
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34 9 18.0 10 1 US-09-6 35 9 18.0 10 1 PCT-USS 37 8.4 16.8 10 1 US-08-3 38 8.4 16.8 10 1 US-08-4 39 8.4 16.8 10 1 US-08-4 40 8.4 16.8 10 1 US-08-4 41 8.4 16.8 10 1 US-08-4 42 8.4 16.8 10 1 US-08-4 43 8.4 16.8 10 1 US-08-4 44 8.4 16.8 10 1 US-08-4 45 8.4 16.8 10 1 US-09-6 46 8.4 16.8 10 1 US-09-6 47 8.4 16.8 10 1 US-09-6 48 8.4 16.8 10 1 US-09-6 49 8.4 16.8 10 1 US-09-6 40 8.4 16.8 10 1 US-09-6 41 8.4 16.8 10 1 US-09-6 42 8.4 16.8 10 1 US-09-6 43 8.4 16.8 10 1 US-09-6 44 8.4 16.8 10 1 US-09-6 45 8.4 16.8 10 1 US-09-6 46 8.4 16.8 10 1 US-09-6 47 10 1 US-09-6 48 8.4 16.8 10 1 US-09-6 49 8.4 16.8 10 1 US-09-6 40 8.4 16.8 10 1 US-09-6 41		C			
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Sequence 50 Sequence 4, Sequence 1, Sequence 10 Sequence 50 Sequence 44 Sequence 44 Sequence 44 Sequence 16 Sequence 16 Sequence 16 Sequence 21 Years 21 Length 17;	nσ	114 17A: 1 60/ 2 60/ 2 60/ 2 10/ 2 10/	CONDI CONDI OF UP GROWI B502 388: £ Lyon Fifth 00 1 1 1 1 1 1 1 1 1 1 1 1 1	cion Us camela en, Jan omb, Jan METHC	
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Sequence 50 Sequence 4, Sequence 1, Sequence 10 Sequence 10 Sequence 54 Sequence 44 Sequence 44 Sequence 16 Sequence 16 Sequence 16 Sequence 21 Sequence 27 Length 17;	13.8; DB	064	RELATED TO LEVE ENDOTHELIAL OR	LLIGNMENTS 1040 REAGENT FOR THE	99-825-574-50 US91-01822A-4 US91-02628-4 B-91-02628-4 B-308-894-6 B-308-474A-3 B-388-353-482 B-388-353-585 B-488-551B-442 B-488-551B-442 B-488-551B-442 B-488-551B-442 B-488-551B-442
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					54444 1168 1168 1168

Best Local Similarity 76.9 Matches 13; Conservative

76.5%;

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Pred. No. 1; 2; Mismatches

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Indels

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RESULT 3
US-08-292-620A-553/c
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; ORGANISM: Mus sp.
US-09-371-772B-3813
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn version 3.0 SEQ ID NO 3813 LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 553, Application US/08292620A Patent No. 5837542
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions ReTITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor FILE REFERENCE: MBHB00,876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 68/584,040
PRIOR PILING DATE: 1995-01-08
PRIOR FILING DATE: 1996-01-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            atent No. 656612
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Pavco, Pam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 14225
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                                                                                                                                                                                                                                                           TITLE OF INVENTION:
            COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: Btorage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C.
                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                  PPLICANT:
                                                                                                                                                                         STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                          CITY:
                                                                                                    COUNTRY: U.S.A. ZIP: 90071-2066
                                                                                                                                                                                                             ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2072 TGAGCCATCCAAAGAAA 2088
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                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION:
                                                                                                                                      ET: Suite 4700
: Los Angeles
S: California
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F INVENTION: RII
F INVENTION: DI
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Stinchcomb, Dan
Escobedo, Jaime
                                                                                                                                                                                         E: Lyon & Lyon
633 West Fifth Street
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Dan T. Stinchcom
James McSwiggen
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Word Perfect 5.1
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RIBOZYME TREATMENT OF
DISEASES OR CONDITIONS
RELATED TO LEVELS OF
INTRACELLULAR ADHESION
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                   IBM P.C. DOS 5.0
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Pred. No. 1;
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                                                                                                                              US-09-071-845-553/c
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FILING DATE: January 19, 1993
FILING DATE: January 19, 1993
FILING DATE: January 19, 1993
                                                                                           Sequence 553, Application US/09071845 Patent No. 6132967
                                                                                                                                                                                                                                                        Matches 12; Conservative
                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 32,327
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: December 7, ATTORNEY/AGENT INFORMATION:
             APPLICANT:
                              APPLICANT:
                                             APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 15 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Warburg, Richard
REGISTRATION NUMBER: 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/292,620A
FILING DATE: August 17, 1994
                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                           2058 CAGAGCAAAAGA 2069
                                                                                                                                                                                             14 CAGAGCAAAAGA 3
                                                             Susan Grimm
                                                                                                                                                                                                                                                                           24.0%;
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Pred. No.
                                                                                                                                                                                                                                                            Mismatches
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COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                      PRIOR APPLICATION DATA:
                                                                                                                             OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
 APPLICATION NUMBER: FILING DATE: August APPLICATION NUMBER:
                                                                       FILING DATE:
CLASSIFICATION:
                                                                                                          APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                          CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Dan T. Stinchcomb
James McSwiggen
Sean Sullivan
Kenneth G. Draper
VENTION: RIBOZYME TREATMENT OF
                                                                                                                                                                               IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE-1 (I-CAM-1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DISEASES OR CONDITIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                       RELATED TO
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17, 1994
08/008,895
                                     US/08/292,620
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                                                                                                                                                                  DOS 5.0
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TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:

553:

LENGTH: 15 base pairs TYPE: nucleic acid STRANDEDNESS: single

POLOGY:

linear

Score 12; Pred. No.

DB 1; 2.8;

Length 15;

0

REFERENCE/DOCKET NUMBER: 20
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600

208/149

TELEPHONE: (213) 955-0440

ATTORNEY/AGENT INFORMATION:

FILING DATE: Janua: APPLICATION NUMBER: FILING DATE: December

JMBER: 07/989,849 December 7, 1992

January 19, 1993

NAME: Warburg, Richard J REGISTRATION NUMBER: 32,

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US-08-388-353-415/c
; MOLECULE TYPE: US-08-388-353-415
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Best Local Similarity
Matches 12; Conservat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 415, Application US/08388353 Patent No. 6010895
                                                                                                                                     TELEPHONE: (516) 742-434
TELEFAX: (516) 742-436
TELEX: 230 901 SANS UR
INFORMATION FOR SEQ ID NO: 4
                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Digiglio, Frank S.
REGISTRATION NUMBER: 31,346
REFERENCE/DOCKET NUMBER: 9606
TELECOMMUNICATION INFORMATION:
TELEPHONE: (516) 742-4343
                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
TILING DATE: 14-FEB-1995
CLASSIFICATION: 424
                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Deacon, Nicholas J.
                                                        STRANDEDNESS:
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                                                                            TYPE: nucleic acid
                                          TOPOLOGY:
                                                                                                 LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TLE OF INVENTION: NON-PATHOGENIC STRAINS OF HIV-1 MBER OF SEQUENCES: 800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2058 CAGAGCAAAAGA 2069
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New York
f: United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    E: Scully, Scott, Murphy & Presser
400 Garden City Plaza
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Learmont, Jennit
McPhee, Dale A.
Crowe, Suzanne
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                    DNA (genomic)
                                                        single
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US-08-488-551B-415/c

Sequence 415, Application US/08488551B

Patent No. 6015661
GENERAL INFORMATION:

APPLICANT: Nicholas J. Deacon
APPLICANT: Dale A. McPhee
APPLICANT: David Cooper
APPLICANT: David Cooper
TITLE OF INVENTION: NON-PATHOGENIC STRAINS OF HIV-1

TUMBER OF SEQUENCES: 841

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Query Match
Best Local Similarity
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RESULT 7
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Matches 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 702, Application US/08388353
                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/388,353
FILING DATE: 14-FEB-1995
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: DIGIGILIO, Frank S.
REGISTRATION NUMBER: 31,346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                         TELEX: 230 901 SANS UR INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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APPLICANT:
                                                                                                                                                                                               MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 96
TELECOMMUNICATION INFORMATION:
TELEPHONE: (516) 742-4343
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Cooper, David
TITLE OF INVENTION: NON-PATHOGENIC STRAINS OF HIV-1
NUMBER OF SEQUENCES: 800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                             STRANDEDNESS:
                                                                                                                                                                                                                                            TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Scully, Scott, Murphy & Presser STREET: 400 Garden City Plaza CITY: Garden City
                                                                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                              TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                 ENGTH:
                                                                              2094 AGCTCTCTGG 2103
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                                                                                                                                                                                                                                                                 10 base pairs
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(516) 742-4366
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                                                                                                               Conservative
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                                                                                                                                                                                                                 linear
                                                                                                                                                                                              DNA (genomic)
                                                                                                                                                                                                                           single
                                                                                                               20.0%; Score 10; DB 1; 100.0%; Pred. No. 11; tive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                  9606
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                                                                                                                                               Length 10
                                                                                                                 Indels
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                                                                                                                 Gaps
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ADDRESSEE: SCULLY, S

ADDRESSEE:

STREET: 400 CITY CITY CARDEN CITY

400 GARDEN CITY

SCOTT, MURPHY & PRESSER

STATE: NEW YORK

COUNTRY: U.S.A. ZIP: 11530-0299

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                                                                                                                                                                                                                                                                                                                                                                           Sequence 702, Application US/08488551B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 960
TELECOMMUNICATION INFORMATION:
TELEPHONE: (516) 742-4343
TELEPHONE: (516) 742-4366
INFORMATION FOR SEQ ID NO: 415:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                     tent No.
                                                                                                                                                                                                                                                                              APPLICANT:
APPLICANT:
       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                       APPLICANT: Dale A. McPhee
APPLICANT: David Cooper
TITLE OF INVENTION: NON-PATHOGENIC STRAINS OF HIV-1
NUMBER OF SEQUENCES: 841
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 17-MAY-1995
ATTORNEY/AGENT INFORMATION:
NAME: FRANK S. DIGIGLIO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                      ADDRESSEE: SCULLY, S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 08/388,353
FILING DATE: 14-FEB-1995
APPLICATION NUMBER: PN3021/95
FILING DATE: 17-MAY-1995
                                                                                                          ZIP: 11530-0299
                                                                                                                                                                 CITY: GARDEN CITY
                                                                                                                                                                                    STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TILING DATE: 14-FEB-1994
ARPLICATION NUMBER: PM4002 (AU)
SILING DATE: 21-FEB-1994
ARPLICATION NUMBER: PN0284 (AU)
SILING DATE: 23-DEC-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 h 20.0%; Score 10; DB 1; Length 10; Similarity 100.0%; Pred. No. 11;
                                                                                                                                                 NEW YORK
                                                                                                                                                                                  E: SCULLY, SCOTT, MURPHY & PRESSER 400 GARDEN CITY PLAZA
                                                                                                                                                                                                                                                                                                                Nicholas J. Deacon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10 base pairs
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                                                                                                                               U.S.A.
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MMBER: PN0284 (AU)
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RESULT 10 US-08-173-489C-274

Sequence 274, Application US/08173489C

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; LENGTH: 11
; TYPE: DNA
; ORGANISM: Human
US-09-657-013-94
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US-08-488-551B-702
                                                                                                                                                                SEQ ID NO 94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                Matches
                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 9
Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                            NUMBER OF SEQ ID NOS:
SOFTWARE: PatentIn ver
                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/657,013
CURRENT FILING DATE: 2000-09-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                09-657-013-94
                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/152,778
PRIOR FILING DATE: 1999-09-07
                                                                                                                                                                                                                                                                          APPLICANT: Van den Veyver, Ignatia B
APPLICANT: Amir, Ruthie
APPLICANT: Francke, Uta
APPLICANT: Francke, Uta
TITLE OF INVENTION: Methods of Identifying Mutations in a
TITLE OF INVENTION: Containing Gene or Protein in Neurode
FILE REFERENCE: HO-P01893US1/0995371
                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Zoghbi, Huda Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (516) 742-4366 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER:
FILING DATE: 17-MAN
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2059 AGAGCAAAAG 2068
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o. 670981
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EDNESS: single
                                                                                                                                                                             PatentIn version 3.1
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                                Conservative
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                                              20.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20.0%; Score 10;
100.0%; Pred. No.
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                                                Score 10; DB 1;
Pred. No. 9.9;
                                                             Length 11;
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                                  Indels
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                                Gaps
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Patent No. 5861244

GENERAL INFORMATION:

HEPBURN, A. G.

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COMPUTER: IBM PC/XT/AT
COMPUTER: IBM PC/XT/AT
OPERATING SYSTEM: MS-DOS version 6.2
SOFTWARE: Wordperfect Version 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/173,489C
FILING DATE: 22 DEC 1993
CLASSIFICATION DATA:
APPLICATION UNDER: US 07/968,436
FILING DATE: 29 OCT 1992
APPLICATION NUMBER: US 07/968,436
FILING DATE: 29 OCT 1992
APPLICATION INFORMATION:
NAME: Handelman, Joseph H.
REGISTRATION NUMBER: 26,179
REFERENCE/DOCKET NUMBER: U9518-6
TELECOMMUNICATION INFORMATION:
TELEPAX: (attorney) (212) 708-1880
TELEPAX: (attorney) (212) 246-8959
INFORMATION FOR SEQ ID NO: 274:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 bases
                                                                                                    GENERAL INFORMATION:
APPLICANT: DUTTA, SUKANTA K.
APPLICANT: BISMAS, Biswajit
APPLICANT: VEMULAPALLI, Ramesh
TITLE OF INVENTION: A SIZE-VARIABLE STRAIN-SPECIFIC PROTECTIVE ANTIGEN FOR
TITLE OF INVENTION: POTOMAC HORSE FEVER
                                                                                                                                                                                                                               9-157-257-17
Sequence 17, Application US/09157257
Patent No. 6375954
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
FILE REFERENCE: 8172-9016
CURRENT APPLICATION NUMBER: US/09/157,257
CURRENT FILING DATE: 1998-09-18
EARLIER APPLICATION NUMBER: 60/059,252
EARLIER FILING DATE: 1997-09-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ANTI-SENSE: no
PUBLICATION INFORMATION:
RELEVANT RESIDUES IN SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HYPOTHETICAL: yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: PROFILE DIAGNOSTIC SCIENCES, INC.
STREET: 510 EAST 73RD STREET,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: other nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DESCRIPTION: third strand derived from DESCRIPTION: 168 region in Seq ID No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS: single stranded
                                                                                                                                                                                                                                                                                                                                                                                                             2080 CCAAAGAAACA 2090
                                                                                                                                                                                                                                                                                                                                                                                                                                                         10;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
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5861244273
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US-09-249-155A-189
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 12
US-09-157-257-32
US-09-249-155A-189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/157,257
CURRENT FILING DATE: 1998-09-18
EARLIER APPLICATION NUMBER: 60/059,252
EARLIER FILING DATE: 1997-09-18
NUMBER OF SEQ ID NOS: 48
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 32
SEQ ID NO 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: DUTTA, SUKANTA K.
APPLICANT: BISWAS, Biswajit
APPLICANT: VEMULAPALLI, Ramesh
TITLE OF INVENTION: A SIZE-VARIABLE STRAIN-SPECIFIC PROTECTIVE ANTIGEN FOR
TITLE OF INVENTION: POTOMAC HORSE FEVER
FILE REFERENCE: 8172-9016
                                                 NUMBER OF SEQ ID NOS: 346
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 189
LENGTH: 11
                                                                                                                                                                                                                                                                                                                                          Sequence 189, Application US/09249155A Patent No. 6538173 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
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Patent No. 6375954
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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Best Local Similarity
                                                                                                                                                                                                                                                          TITLE OF INVENTION: Compositions and Methods for Wound TITLE OF INVENTION: Compositions and Methods for Wound FILE REFERENCE: 00486.78502 CURRENT ADDITION:
                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/249,155A CURRENT FILING DATE: 1999-02-12 PRIOR APPLICATION NUMBER: US 60/074,737 PRIOR FILING DATE: 1998-02-13 PRIOR APPLICATION NUMBER: US 60/097,937 PRIOR FILING DATE: 1998-08-26 PRIOR APPLICATION NUMBER: US 60/102,051
                                                                                                                          PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 1998-
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                 ORGANISM: Mus musculus
                                  TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH:
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Pred. No. 13;
0; Mismatches
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Pred. No. 13;
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US-07-724-500B-4
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US-08-461-418B-4
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US-07-724-500B-4
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                                                                      Sequence 4, Application US/08461418B Patent No. 5874564
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                                           GENERAL INFORMATION:
APPLICANT: Ecker
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APPLICANT: Ecker
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MEDIUM TYPE: DISKETT
COMPUTER: IBM PS/2
OPERATING SYSTEM: PC
           TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: IS TELECOMMUNICATION INFORMATION: TELEPHONE: (215) 568-3100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 2
                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: PCT/US91/01822
FILING DATE: 19 MARTON 1991
NTORNBY AGENT INFORMATION:
NAME: John W. Caldwell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: WORDPERFECT 5.0
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                      18.0%;
Local Similarity 66.7%;
les 6; Conservarino
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FILING DATE: June 27, 1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Woodcock Washburn Kurtz
ADDRESSEE: Mackiewicz & No. 5736294ris
STREET: One Liberty Place - 46th Floor
                                                                                                                                                                                                                                                                                                                                                                                                      ENGTH:
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                                                                                                                                                                                                 2095 GCTCTCTGG 2103
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10; Conservative
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                                          Ecker et al.
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Expression Through RNA Mimicry
                           Reagents And Methods For Modulating Gene
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Pred. No. 17;
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Pred. No. 1
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Patent No. 6
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Best Local Similarity 66.7%;
Matches 6; Conservative
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: FORDS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UNMERR: US/08/388,353
FILING DATE: 14-FEB-1995
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ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz & No. STREET: One Liberty Place - 46th floor
                                                                                                                                                                                                                                                                                       APPLICANT: Crowe, Suzanne
APPLICANT: Cooper, David
TITLE OF INVESTION: NON-PATHOGENIC STRAINS OF HIV-1
NUMBER OF SEQUENCES: 800
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REFERENCE/DOCKET NUMBER: IS
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 16-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Paul K. Legaard
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MEDIUM TYPE: 3.5 inc
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PRIOR APPLICATION DATA:
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CURRENT APPLICATION DATA:
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CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                        STATE: New York
                                                                                                                                                                                                                        CITY: Garden City
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STRANDEDNESS: single
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OPERATING SYSTEM: PC-DOS/MS-DOS
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Pred. No.
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; TOPOLOGY: linear; MOLECULE TYPE: DNA (genomic) US-08-388-353-32
                                                                   ; TOPOLOGY: 1:
; MOLECULE TYPE:
US-08-388-353-33
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Best Local Similarity 100.0%;
Matches 9; Conservative
   Query Match
Best Local Similarity
Matches 9; Conserv
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Patent No. 6010895
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (516) 742-4343
TELEFAX: (516) 742-4366
TELEX: 230 901 SANS UR
INFORMATION FOR SEQ ID NO: 32:
                                                                                                                                                                                    TELEX: 230 901 SANS U
                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Diciglio, Frank S.
REGISTRATION NUMBER: 31,346
REFERENCE/DOCKET NUMBER: 960
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/388,353
FILING DATE: 14-FEB-1995
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MEDIUM TYPE: Floppy
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TITLE OF INVENTION: NON-PATHOGENIC STRAINS OF HIV-1
TUMBER OF SEQUENCES: 800
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                                                                                                               STRANDEDNESS:
                                                                                                                             TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                   (516) 742-4366
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18.0%; Score 9; DB 1;
100.0%; Pred. No. 17;
tive 0; Mismatches
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Mismatches
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Query Match
Best Local Similarity
Watches 9; Conserve
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US-08-388-353-416/c
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US-08-388-353-414
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US-08-388-353-414/c
                                                                                                                                                                   Sequence 416, Application US/08388353 Patent No. 6010895
                                                                                                                                                                 Patent No.
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                                                                                                                 GENERAL INFORMATION:
APPLICANT: Deacon,
APPLICANT: Learmon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEX: 230 901 SANS UR INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PATEN: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/388,353
FILING DATE: 14-FEB-1995
CLASSIFICATION: 424
                             APPLICANT: Cooper, David
TITLE OF INVENTION: NON-PATHOGENIC STRAINS OF HIV-1
NUMBER OF SEQUENCES: 800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION: TELEPHONE: (516) 742-4343
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: DiGiglio, Frank S.
REGISTRATION NUMBER: 31,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Cooper, David
TITLE OF INVENTION: NON-PATHOGENIC STRAINS OF HIV-1
                                                                                  APPLICANT:
                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
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STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET:
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400 Garden City Plaza
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Crowe, Suzanne
                                                                                McPhee, Dale A. Crowe, Suzanne
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (516) 742-4366
                                                                                                               Learmont, Jennifer C.
                                                                                                                                Deacon, Nicholas J.
                                                                                                                                                                                                                                                                                                                                  Conservative
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Scully, Scott, Murphy & Presser
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100.0%; Pred. N
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COUNTRY:

United States

Garden City New York

400 Garden City Plaza

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RESULT 20
US-08-388-353-701
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REGISTRATION NUMBER: 31,346
REFERENCE/DOCKET NUMBER: 9606
TELECOMMUNICATION INFORWATION:
TELEPHONE: (516) 742-4343
TELEPHONE: (516) 742-4343
TELEPHONE: (516) 742-4366
TELEZ: 230 901 SANS UR
INFORMATION FOR SEQ ID NO: 416:
SEQUENCE CHARACTERISTICS:
COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/388,353
FILING DATE: 14-FEB-1995
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: DiGiglio, Frank S.
REGISTRATION NUMBER: 31,346
REFERENCE/DOCKET NUMBER: 9606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequence 701,
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
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FILING DATE: 14-FEB-1995
CLASSIFICATION: 424
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les 9; Conserv
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                  CITY: Garden City
STATE: New York
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400 Garden City Plaza
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                                                                                                                                                                                                                                                                 United States
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Crowe, Suzanne
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100.0%; Pred. No.
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                                            Query Match
Best Local Similarity
Thes 9; Conserv
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INFORMATION FOR SEQ ID NO:
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LENGTH: 10 base pairs
                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/388,353
FILING DATE: 14-FEB-1995
FILING DATE: 14-FEB-1995
CLASSIFICATION: 424
ANDE: DISCRIPT INFORMATION:
NAME: DISCRIPT SEPAPLOR
                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 96
TELECOMMUNICATION INFORMATION:
TELEPHONE: (516) 742-4343
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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CORRESPONDENCE ADDRESS:
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APPLICANT: Cooper, DON'DATHOGENIC STRAINS OF HIV-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                  TYPE: nucleic STRANDEDNESS:
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ses 9; Conserv
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STRANDEDNESS: single
                                                                                                                                                      TOPOLOGY:
                                                                                                                                                                                                                                                               TELEFAX: (516) 742-4366
                                                                                                                                                                                                                                                                            TELEPHONE:
                                                                                                                                                                                                                                                                                                                          NAME: DiGiglio, Frank S. REGISTRATION NUMBER: 31,
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                            2095 GCTCTCTGG 2103
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                                                                                                                                                                              10 base pairs
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Crowe, Suzanne
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                                                         Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA (genomic)
                                                                                                                                   DNA (genomic)
                                                                                                                                                                   single
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100.0%; Pred. No.
                                                                         18.0%; Score 9;
100.0%; Pred. N
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Mismatches
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                                                            Indels
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US-08-488-551B-33/c
; Sequence 33, Application US/08488551B
; Patent No. 6015661
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                                                                                                                                                          RESULT 23
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                                                                                                                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                                                                                              Query Match
               GENERAL INFORMATION:
APPLICANT: Nicholas J. Deacon
APPLICANT: Dale A. McPhee
APPLICANT: David Cooper
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: PM3864 (AU)
FILING DATE: 14-FEB-1994
APPLICATION NUMBER: PM4002 (AU)
FILING DATE: 21-FEB-1994
APPLICATION NUMBER: PM20284 (AU)
FILING DATE: 23-DEC-1994
APPLICATION NUMBER: US 08/388,35
FILING DATE: 14-FEB-1995
APPLICATION NUMBER: PM3021/95
APPLICATION NUMBER: PM3021/95
APPLICATION NUMBER: PM3021/95
APPLICATION NUMBER: PM3021/95
ATTORNEY/AGENT INFORMATION:
NAME: FRANK S. DIGIGLIO
REFERENCE/DOCKET NUMBER: 9606Z
TELECOMMUNICATION INFORMATION:
TELEPHONE: (516) 742-4366
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (516) 742-430
[NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 1
FILING DATE: 07-JUN-
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: 11 MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Dale A. McPhee
APPLICANT: David Cooper
TITLE OF INVENTION: NON-PATHOGENIC STRAINS OF HIV-1
NUMBER OF SEQUENCES: 841
     TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
                                                                                                                                                                                                                                                                                      Local Similarity les 9; Conserv
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OPERATING SYSTEM: PC-DOS/MS-DOS
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400 GARDEN CITY PLAZA
                                                                                                                                                                                                                                                                                                                                                                                                                                                               10 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      742-4366
                                                                                                                                                                                                                                                                                      18.0%; Score 9; DB 1;
100.0%; Pred. No. 17;
tive 0; Mismatches
NON-PATHOGENIC STRAINS OF HIV-1
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                                                                                                                                                                                                                                                                                                                          Length 10;
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Best Local Similarity
Watches 9; Conserve
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                                                                                                                                                                                                                                                                                                                                                                        Patent No.
                                                                                                                                                                                                                                                                                                                                                                                        Sequence 414, Application US/08488551B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (516) 742-4366
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
LENGTH: 10 base pairs
                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                             APPLICANT: Nicholas J. Deacon
APPLICANT: Dale A. McPhee
APPLICANT: David Cooper
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (516) 742-4343
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ATTORNEY/AGENT INFORMATION: NAME: FRANK S. DIGIGLIO
                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
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                                                                                                                                                                CITY: GARDEN CITY STATE: NEW YORK
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STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: PN0284 (AU) FILING DATE: 23-DEC-1994 APPLICATION NUMBER: US 08/388,353
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                                                                                                                                              COUNTRY:
                                                                                                                                                                                                      STREET:
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FILING DATE: 14-FEB-1995
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                                                                                                                                                                                                                      ADDRESSEE:
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                                                                                                                               11530-0299
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                                                                                                                                                                                                                                                          David Cooper
David Cooper
VENTION: NON-PATHOGENIC STRAINS OF HIV-1
                                                                                                                                                U.S.A.
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100.0%; Pred. No.
Live 0; Mismatc
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US/08/488,551B
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FILING DATE: 07-JUN-LIOR APPLICATION DATA: APPLICATION NUMBER:

07-JUN-1995

14-FEB-1994

PM4002 (AU) PM3864 (AU)

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RESULT 25
US-08-488-551B-416/c
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FILING DATE: 14-FEB-1995
APPLICATION NUMBER: PN3021/95
APPLICATION NUMBER: PN3021/95
FILING DATE: 17-MAY-1995
ATTORNEY/AGENT INFORMATION:
NAME: FRANK S. DIGIGIO
REFERENCE/DOCKET NUMBER: 9606
TELECOMMUNICATION INFORMATION:
TELEPHONE: (516) 742-4343
TELEPHONE: (516) 742-4343
TELEPAX: (516) 742-4346
INFORMATION FOR SEQ ID NO: 414:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 base pairs
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                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: U$5/08/488,551B
FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PM3864 (AU)
FILING DATE: 14-FEB-1994
APPLICATION NUMBER: PM4002 (AU)
APPLICATION NUMBER: PM4002 (AU)
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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ADDRESSEE: SCULLY, SCOTT, MURPHY & PRESSER
STREET: 400 GANDEN CITY PLAZA
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APPLICANT: Dale A. McPhee
APPLICANT: David Cooper
TITLE OF INVENTION: NON-PATHO
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APPLICATION NUMBER: PN0284 (AU)
FILING DATE: 23-DEC-1994
FILING DATE: US 08/388,353
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                                   FILING DATE: 21-FEB-1994
APPLICATION NUMBER: PNO284 (AU)
FILING DATE: 23-DEC-1994
APPLICATION NUMBER: US 08/388,3
FILING DATE: 14-FEB-1995
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FILING DATE: 21-FEB-1994
APPLICATION NUMBER: PN3021/95 FILING DATE: 17-MAY-1995
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Query Match
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; MOLECULE TYPE:
US-08-488-551B-416
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                                                                                                                                   TELEFAX: (516) 742-4366
INFORMATION FOR SEQ ID NO: 7
SEQUENCE CHARACTERISTICS:
LENGTH: 10 base pairs
                                                                                                                                                                                                                                                                                           FILING DATE: 17-MAY-1995
ATTORNEY/AGENT INFORMATION:
NAME: FRANK S. DIGIGLIO
REFERENCE/DOCKET NUMBER: 9606
TELECOMMUNICATION INFORMATION:
TELEPHONE: (516) 742-4343
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 961
TELECOMMUNICATION INFORMATION:
TELEPHONE: (516) 742-4343
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MOLECULE TYPE:
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                                                                                                       LENGTH: 10 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 21-FEB-1994
APPLICATION NUMBER: PN0284 (AU)
FILING DATE: 23-DEC-1994
APPLICATION NUMBER: US 08/388,353
FILING DATE: 14-FEB-1995
APPLICATION NUMBER: PN3021/95
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                               STRANDEDNESS: single
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FILING DATE:
PILING DATE:
21-FEB-1994
PILICATION NUMBER:
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o. 6015661
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18.0%; Score 9; DB 1; milarity 100.0%; Pred. No. 17; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: SCULLY, SCOTT, MURPHY & PRESSER 400 GARDEN CITY PLAZA
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RESULT 27
US-08-488-551B-703
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Best Local Similarity
Thehes 9; Conserve
RESULT 28
US-09-034-205-50/c
; Sequence 50, Application US/09034205
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                                                                                                                                                                                                                                   ; MOLECULE TYPE: DNA US-08-488-5518-703
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Best Local Similarity
""" Conser
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Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: U.S.A.

ZIP: 11530-0299
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Vers
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/488,551B
FILING DATE: 07-JUN-1995
PRIOR APPLICATION NUMBER: PM3864 (AU)
APPLICATION NUMBER: PM3864 (AU)
                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 14-FEB-1994
APPLICATION NUMBER: PM4002 (AU)
FILING DATE: 21-FEB-1994
APPLICATION NUMBER: PN0284 (AU)
FILING DATE: 23-DEC-1994
APPLICATION NUMBER: US 08/388,353
FILING DATE: 14-FEB-1995
APPLICATION NUMBER: PN3021/95
FILING DATE: 17-MAY-1995
APPLICATION TORNATION:
NAME: PENNY S DITTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (516) 742-4366 INFORMATION FOR SEQ ID NO: '
                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: NON-PATHOGENIC STRAINS OF HIV-1
NUMBER OF SEQUENCES: 841
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 96
TELECOMMUNICATION INFORMATION: (516) 742-4343
                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                    TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: GARDEN CITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: FRANK S. DIGIGLIO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                      LENGTH:
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                                                                                              GCTCTCTGG 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Application US/084885518
                                                                                                                                                                                                                                                                                                                     10 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            David Cooper
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dale A. McPhee
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                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                single
                                                                                                                                                            18.0%; Score 9; DB 1;
100.0%; Pred. No. 17;
tive 0; Mismatches
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/09/034,205
FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: MACKHIGHT, Kamrin T.
REGISTRATION UNMBER: 38,230
REFERENCE/DOCKET NUMBER: FORS-03268
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 397-8338
INFORMATION FOR SEQ ID NO: 50:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 base pairs
TYPE: nucleic acid
TTPE: nucleic acid
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US-08-934-097A-50/c
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GENERAL INFORMATION:
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APPLICANT:
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MOLECULE TYPE:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: MEDLEN & CARROLL, LLP
STREET: 220 Montgomery Street, Suite 2200
COUNTRY: USA
ZIP: 94104
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                  TITLE OF INVENTION: Polymorphism Analysis By Nucleic Acid TITLE OF INVENTION: Structure Probing With Structure-Bridging TITLE OF INVENTION: Oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Fore, Lance
APPLICANT: Neri, Bruce P.
TITLE OF INVENTION: TARGET-DEPENDENT REACTIONS USING
TITLE OF INVENTION: STRUCTURE-BRIDGING OLIGONUCLEOTIDES
                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: MEDLEN & CARROLL, LLP
STREET: 220 Montgomery Street, Suite 2200
                                                                                                                                                                                                                                                                                                           APPLICANT: Lyamichev, Victor I. APPLICANT: Brow, Mary Ann D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Lyamichev, Victor APPLICANT: Brow, Mary Ann D.
                                                                                                       CITY: San Francisco
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DESCRIPTION:
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Neri, Bruce P.
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/desc = "DNA"
Floppy disk
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RESULT 30
US-09-677-218B-50/c
; Sequence 50, Application US/09677218B
; Patent NO. 6355437
; GENERAL INFORMATION:
; APPLICANT: Lyamichev, Victor I.
Brow, Mary Ann D.
Tance
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (415) 397-83:
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: MacKnight, Kamrin T.
REGISTRATION NUMBER: 38,230
REFERENCE/DOCKET NUMBER: FORS-02980
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 705-8410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA RELEASE #1.0, Version #1.30
CURRENT APPLICATION DATA:
                   INFORMATION
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                                                                                                                                                                                                                  COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PA-EDICATION DATA:
SOFTWARE: PATENTIA Release #1.0, Vers
APPLICATION DATA:
APPLICATION UNMER: US/09/677,218B
FILING DATE: 02-Oct-2000
CLASSIFICATION: CUNKNOWN>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
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                                                                      APPLICATION NUMBER: 09/034,205
FILING DATE: «Unknown»
ATTORNBY/AGENT INFORMATION:
NAME: MacKnight, Kamrin T.
REGISTRATION NUMBER: 38,330
REFERENCE/DOCKET NUMBER: FORS-03268
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: TARGET-DEPENDENT REACTIONS USING
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10 GCTCTCTGG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  h 18.0%; Score 9; DB:
Similarity 100.0%; Pred. No. 17
9; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: MEDLEN & CARROLL, LLP
STREET: 220 Montgomery Street, Suite 2200
CITY: San Francisco
STATE: CA
              TELEPHONE: (415) 705-8410
TELEFAX: (415) 397-8338
N FOR SEQ ID NO: 50:
                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: USA
  CHARACTERISTICS
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5. 17;
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DESCRIPTION: /deec = "DLA"

SEQUENCE DESCRIPTION: SEQ ID NO: 50:
US-09-677-218B-50
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                                                                                                                                                                                                                                                                                                                                      RESULT 32
                                                                                                                                                                                                                                                                Sequence 4, Application US/09255899
Patent No. 6368863
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 50, Application US/09677192 Patent No. 6358691
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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Best Local S
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CURRENT APPLICATION NUMBER: US/09/677,192
CURRENT FILING DATE: 2000-10-02
PRIOR APPLICATION NUMBER: 09/034,205
PRIOR FILING DATE: 1998-03-03
PRIOR FILING DATE: 1998-03-03
NUMBER OF SEQ ID NOS: 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: TAKGET TITLE OF INVENTION: OLIGONUCLEOTIDES
--- PERFERENCE: FORS-04708
US/09/677
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Description of Artificial Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 10
ZIP: 19103
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch disk, 1.44 Mb
COMPUTER: IBM PC compatible
COMPUTER: FIRM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2095 GCTCTCTGG 2103
                                                                                                                                 NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESSE: Woodcock Washburn Kurtz Mackiewicz & No. 6368863ris LLP
STREET: One Liberty Place - 46th Floor
CITY: Philadelphia
                                                                                                                                                                                                          APPLICANT: Ecker et al.
APPLICANT: Ecker et al.
TITLE OF INVENTION: Reagents And Methods For Modulating Gene
Expression Through RNA Mimicry
                                                                                                                                                                                                                                                                                                                                                                                                                      2095 GCTCTCTGG 2103
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9; Conserv
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Brow, Mary Ann D.
Fors, Lance
                                                                                                                    STATE: PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 10 base pairs
                                                                                                     COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18.0%; Score 9; ilarity 100.0%; Pred. No. Conservative 0; Mismatcl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
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APPLICANT: Anderson, Todd
APPLICANT: Dahlberg, James
ITITLE OF INVENTION: Target-Dependent Reactions Using Structure-Bridging Oligonucleoti
FILE REFERENCE: FORS-04012
CURRENT APPLICATION NUMBER: US/09/402,618B
CURRENT FILING DATE: 2000-07-18
PRIOR APPLICATION NUMBER: PCT/US98/03194
PRIOR FILING DATE: 1998-05-05
NUMBER OF SEQ ID NOS: 128
SOFTWARE: Patentin version 3.0
SEQ ID NO 50
LENGTH: 10
                                                                                                                                                                                               ; TYPE: DNA
GRANISM: Artificial Sequence
; PEATURE:
; OTHER INFORMATION: Synthetic
US-09-402-618B-50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Thes 6; Conserve
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US-09-402-618B-50/c
US-09-825-574-50/c
                 RESULT 34
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APPLICATION NUMBER: 08/461,418

FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:

NAME: Paul K. Legaard

REGISTRATION NUMBER: 38,534

REFERENCE/DOCKET NUMBER: ISIS-1998

TELECOMMUNICATION INFORMATION:

TELEPHONE: 215-568-3100

TELEPHAX: 215-568-3439

INFORMATION FOR SEQ ID NO: 4:

SEQUENCE CHARACTERISTICS:
LENGTH: 10 bases

TYDER: TRACTIC SCIENTING
                                                                                                                                    Matches
                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 50, Appl
Patent No. 670981
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APPLICATION NUMBER: US/09/255,899
FILING DATE: 23-Feb-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2095 GCTCTCTGG 2103
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                                                                                               2095 ecrercies 2103
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                                                                  GCTCTCTGG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: WordPerfect 6.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Application US/09402618B
                                                                                                                                Conservative
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66.7%;
                                                                                                                                                18.0%;
                                                                                                                              Score 9; DB 1;
; Pred. No. 17;
0; Mismatches
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Pred. No. 17;
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US-09-825-574-50
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APPLICATION NUMBER: 08/934,097

FILING DATE: <UNKnown>
ATTORNEY/AGENT INFORMATION:
NAME: MacKnight, Kamzin T.
REGISTRATION NUMBER: 38,230

REFERENCE/DOCKET NUMBER: FORS-02980

TELEPHONE: (415) 705-8410

TELEPHONE: (415) 705-8410

TELEPAX: (415) 397-8338

INFORMATION FOR SEQ ID NO: 50:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 base pairs
TYPE: nucleic acid
STRANDENNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 9; Conserv
                                                                                                                                                                                                                                                                                    Sequence 4, Application PC/TUS9101822A GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patent No. 6709819
GENERAL INFORMATION:
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                                                                                                                                                                                                           APPLICANT: Ecker et al.
TITLE OF INVENTION: REAGENTS AND METHODS FOR MODULATING
TITLE OF INVENTION: GENE EXPRESSION THROUGH RNA MIMICRY
                                                                                                                                                                           NUMBER OF SEQUENCES: S
ZIP: 19103
                                                                  CITY: Philadelphia
STATE: PA
                                                                                                           ADDRESSEE: Woodcock Washburn Kurtz
ADDRESSEE: Mackiewicz & Norris
STREET: One Liberty Place - 46th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/825,574
FILING DATE: 03-Apr-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "DNA"
SEQUENCE DESCRIPTION: SEQ ID NO: 50:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 51
CORRESPONDENCE ADDRESS:
ADDRESSEE: MEDLEN & CARROLL, LLP
STREET: 220 Montgomery Street, SI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Polymorphism Analysis By Nucleic Acid
Structure Probing With Structure-Bridg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Lyamichev, Victor I.
                                                                                                                                                                                                                                                                                                                                                                                                                           10 GCTCTCTGG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: San Francisco
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18.0%; Score 9;
100.0%; Pred. No.
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CLASSIFICATION: 435
PRIOR APPLICATION DATA: 437,090
FILING DATE: MARCH 21,1990
ATTORNEY/AGENT INFORMATION: 1990
ATTORNEY/AGENT INFORMATION: REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISIS-0109
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
TELEPAX: (215) 568-3439
INFORMATION FOR SEQ. ID NO: 4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 4, Application PC/TUS9102628 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 18.0%;
Best Local Similarity 66.7%;
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          -US91-02628-4
                 TELEFAX: (215) 568-3439 INFORMATION FOR SEQ ID NO:
                                               REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: IS:
TELECOMMUNICATION INFORMATION:
TELECHOONE: (215) 568-3100
                                                                                                                                                        APPLICATION NUMBER: PCT/US91/02628
FILING DATE: 19910417
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 497,090
FILING DATE: March 21, 1990
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM: PC-DOS
SOFTWARE: WORDPERFECT 5.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PS/2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
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CURRENT APPLICATION DATA:
CURRENT APPLICATION NIMBER: PC'
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: REAGENTS AND METHODS FOR MODULATING TITLE OF INVENTION: GENE EXPRESSION THROUGH RNA MIMICRY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Woodcock Washburn Kurtz
ADDRESSEE: Mackiewicz & Norris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                     19103
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WORDPERFECT 5.0
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Pred. No.
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. 17;
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Query Match
Best Local Similarity
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US-08-205-507-10
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                                                                                                                                                                                                                               NAME: JOSEPH LUCCI
REGISTRATION NUMBER: 33,307
REFERENCE/DOCKET NUMBER: ISIS
TELECOMMUNICATION INFORMATION:
TELEPHONE: (21.5) 568-3100
TELEPHAX: (21.5) 568-3439
INFORMATION FOR SEQ ID NO: 10:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 18.0%;
Best Local Similarity 66.7%;
Matches 6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 10,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Phillip E
APPLICANT: Bruice
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: PCT/US93/02059
FILING DATE: March 5, 1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 19103
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz & No. 5543507ris STREET: One Liberty Place - 46th Floor
                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PS/2
OPERATING SYSTEM: PC-DOS
SOFTWARE: WORDPERFECT 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS:
TOPOLOGY: unl
                                                                                                                                                     TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH:
                            2096 CTCTCTGGGC 2105
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                                                                                                                                                                                nucleic acid
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DEDNESS: single
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                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Phillip Dan Cook, Muthiah Manoharan, and Thomas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                           16.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Covalently Cross-Linked Oligonucleotides : 17
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                                                                                                                                                                                                                                                                                              ISIS-1304
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Pred. No.
                                                             3; Mismatches
                                                                           Score 8.4; I
Pred. No. 22;
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17;
                                                                                          DB 1; Length 10;
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RESULT 38 US-08-308-894-6

Sequence 6, Application US/08308894 Patent No. 5571672

GENERAL INFORMATION:

Slavicek, James M.

APPLICANT: Garner, Karen J.
APPLICANT: Schreiber, David E.
TITLE OF INVENTION: GYPSY MOTH GENOTYPE ASSAY
NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:

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; TOPOLOGY: linear; MOLECULE TYPE: oligonucleotide US-08-308-894-6
                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-808-474A-3/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 3, Application US/08808474A Patent No. 5856103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: U.S.A.
ZIP: 53705-2398
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, V.
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (608) 231-9508 INFORMATION FOR SEQ ID NO:
                                                               SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Stockhausen, Janet I.
REGISTRATION NUMBER: 34,256
TELECOMMUNICATION INFORMATION: (608) 231-502
                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Locke Purnell Rain Harrell
STREET: 2200 Ross Avenue, Suite 2200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Stockhausen, Janet
              APPLICATION NUMBER: US/08/808,474A
FILING DATE: 03-MAR-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: METHOD FOR SELECTIVELY RANKING SEQUENCES TITLE OF INVENTION: FOR ANTISENSE TARGETING
                                                                                                                                                                                                                                                                                                                                               APPLICANT: Gray, Donald M. APPLICANT: Clark, Chris L.
                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                       STREET: 2200
CITY: Dallas
STATE: Texas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/308,894 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
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DEDNESS: single
DGY: linear
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Mayfield,
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Pred. No. 22,
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                                                                                 Version #1.30
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Query Match
Best Local Similarity
Watches 9; Conserva
Query Match
Best Local Similarity
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                                                                         US-08-388-353-442
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REFERENCE/DOCKET NUMBER: UTDA
TELECOMMUNICATION INFORMATION:
TELEPHONE: (214) 740-8000
TELEPAX: (214) 740-8800
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Patent No.
                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/388
FILING DATE: 14-FEB-1995
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: DiGiglio, Frank S.
REGISTRATION NUMBER: 31,346
REFERENCE/DOCKET NUMBER: 9606
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                            TELEPHONE: (516) 742-436
TELEFAX: (516) 742-4366
TELEX: 230 901 SANS UR
INFORMATION FOR SEQ ID NO: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 10 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Cooper, David
TITLE OF INVENTION: NON-PATHOGENIC STRAINS OF HIV-1
NUMBER OF SEQUENCES: 800
                                                                                      TOPOLOGY: 1
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
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STATE: New York
                                                                                                                  STRANDEDNESS:
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400 Garden City Plaza
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Crowe, Suzanne
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               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linear
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                                                                                      (genomic)
                            16.8%;
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            Score 8.4; DB Pred. No. 22; 0; Mismatches
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Pred. No. :
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                                           DB 1;
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               Indels
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2069 ACTTGAGCCA 2078

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US-08-388-353**-**585/c

Application US/08388353

Patent No.

GENERAL INFORMATION:

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US-08-488-551B-442
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                                                                                                                                                         Patent No.
                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                        equence 442, Application US/08488551B
                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (516) 742-4360
TELEX: 230 901 SANS UR
NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/388,353 FILING DATE: 14-FEB-1995 CLASSIFICATION: 424 CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                              APPLICANT:
                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
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                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                           16.8%;
Local Similarity 90.0%;
les 9; Conservation
                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 10 base pairs
TYPE: nucleic acid (
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: DiGiglio, Frank S. REGISTRATION NUMBER: 31,346 REFERENCE/DOCKET NUMBER: 96
              STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE:
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                             ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
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                                                                                                                                                                                                                                                                                     2094 AGCTCTCTGG 2103
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GARDEN CITY
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             E: SCULLY, SCOTT, MURPHY & PRESSER 400 GARDEN CITY PLAZA
                                                                                                             Dale A. McPhee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           : (516) 742-4343
(516) 742-4366
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Crowe, Suzanne
                                                                                         David Cooper
                                                                                                                           Nicholas J. Deacon
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t, Jennifer C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NON-PATHOGENIC STRAINS OF HIV-1
                                                                             NON-PATHOGENIC STRAINS OF HIV-1
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (516) 742-4366 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 841
CORRESPONDENCE ADDRESS:
ADDRESSE: SCHITT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                            PRIOR APPLICATION DATA:
                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 96
TELECOMMUNICATION INFORMATION:
TELEPHONE: (516) 742-4343
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION: NAME: FRANK S. DIGIGLIO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Nicholas J. Deacon
                                                                                                                                                                                                                               CITY: GARDEN CITY STATE: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 14-FEB-1995
APPLICATION NUMBER: PN30
FILING DATE: 17-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: PN02 FILING DATE: 23-DEC-1994 APPLICATION NUMBER: US 0
APPLICATION NUMBER:
                           APPLICATION NUMBER:
                                                               APPLICATION NUMBER: US/08/488,551B FILING DATE: 07-JUN-1995
                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2069 ACTTGAGCCA 2078
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                                                                                                                                                                                                11530-0299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity 90. 9; Conservative
                                                                                                                                                                                                                                                              E: SCULLY, SCOTT, MURPHY & PRESSER 400 GARDEN CITY PLAZA
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                                                                                                                                                                                                                                                                                                                                                                                                                                Application US/08488551B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA
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MBER: PM4002 (AU)
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MBER: PN0284 (AU)
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             14-FEB-1994
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                             PM3864 (AU)
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PM4002 (AU)
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US 08/388,353

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                                                                                                   RESULT 45
US-09-769-482-16/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-508-753B-166
Sequence 16, Application US/09769482
PATENT NO. 6566130
GENERAL INFORMATION:
APPLICANT: SRIVASTAVA, SHIV
APPLICANT: MOUL, JUDD W.
                                                                                                                                                                                                                                                       Query Match 16.8%;
Best Local Similarity 90.0%;
Matches 9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 472
SEQ ID NO 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 16.8%;
Best Local Similarity 90.0%;
Matches 9; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Yasuhiro FURUICHI
APPLICANT: Yasuhiro FURUICHI
APPLICANT: Yuko SHIBATA
APPLICANT: Hiroko FUNAKI
APPLICANT: Eiji OHARA
APPLICANT: Masanori WATAHIKI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 21-FEB-1994
APPLICATION NUMBER: PN0284 (AU)
FILING DATE: 23-DEC-1994
APPLICATION NUMBER: US 08/388,353
FILING DATE: 14-FEB-1995
APPLICATION NUMBER: PN3021/95
FILING DATE: 17-MAY-1995
ATTORNEY/AGENT INFORMATION:
NAME: FRANK S. DIGIGLIO
REFERENCE/DOCKET NUMBER: 96062
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (516) 742-4366
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/508,753B
CURRENT FILING DATE: 2000-06-16
PRIOR APPLICATION NUMBER: UP 9/270324
PRIOR FILING DATE: 1997-09-18
                                                                                                                                                                                                                                                                                                                                      09-508-753B-166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE: 00162/HG
                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Description of Artificial Sequence: Primer
                                                                                                                                                                                                                                                                                                                                                                                                                                     ENGTH: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION: TELEPHONE: (516) 742-4343
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS: single
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                                                                                                                                                                              TAAGCCATCC 10
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Search completed: November Job time: 1 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 46
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                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/822,250A
CURRENT FILING DATE: 2001-04-02
PRIOR APPLICATION NUMBER: US 08/935,377
PRIOR FILING DATE: 1997-09-22
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PatentIn version 3.2
SEQ ID NO 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21, Applicat Patent No. 6706477 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/179,045
PRIOR FILING DATE: 2000-01-31
NUMBER OF SEQ ID NOS: 67
SOFTWARE: PATENTIN VET. 2.1
SEQ ID NO 16
LENGTH: 10
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Best Local Similarity 90.0%;
Matches 9; Conservative
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APPLICANT: SEGAMA, TAKEHIKO
TITLE (OF INVENTION: PROSTATE-SPECIFIC ANDROGEN-SIGNALING-ASSOCIATED
TITLE OF INVENTION: POYNUCLECTIDE ARRAY
FILE REFERENCE: 04995.0057-00000
CURRENT FILING DATE: 2001-01-26
PRIOR APPLICATION NUMBER: 05/178,772
PRIOR APPLICATION NUMBER: 60/178,772
PRIOR FILING DATE: 2000-01-28
PRIOR FILING DATE: 2000-01-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Zauderer, Maurice
TITLE OF INVENTION: Methods for Producing Polynucleotide Libraries in Vaccinia Virus
FILE REFERENCE: 1821.0010001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                             TYPE: DNA
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                                                                                                                                                                                                                                                                                                                             LENGTH: 10
                                                                                                                2067 AGACTTGAGC 2076
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Pred. No. 22;
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1 20 40.0 20 1 ADG86982  2 2 40.0 20 1 ADG86983	Database: rngdb:*  Database: rngdb:*  Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.  SUMMARIES  Result Query No. Score Match Length DB ID  Description	Geight (c) Ch, us B, 20 B, 20 Caaaag Caaaag NUC O, Ga 1448 1448 1448
C C C C C C C C C C C C C C C C C C C	CC	0 0 000 0 000 0 000 0 0 0 0 0 0 0 0 0
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1 ABH56 191 1 ABH56 191 1 ABH652016 1 ABH62205 1 ABH615694 1 ABH615694 1 ABH696399 1 ABF26457 1 ABF26457 1 ABF296399 1 ABC95308 1 ABC95308 1 ABC95308 1 ABC05733 1 ABF03883 1 ABC05733 1 ABF03882 1 ABF61405 1 ABF161405 1 ABF25416 1 ABF34689 1 ABF34689 1 ABF34689 1 ABF3469 1 ABF36934 1 ABF3688		
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## ALIGNMENTS

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RESULT 1
ADG86982
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                                                                                                                                            conclude shares for an antiberise colligative computation of a nucleix of a nucleix cald controlled colligative activated receptor collection, where the antisense compound inhibits the expression of the pparticle delta and has any of the 6s sequences of 20 amino acids fully defined in the specification. Also included are a compound of 8-80 nucleobases in the specification. Also included are a compound of 8-80 nucleobases in the specification of a preferred target region on a nucleic acid molecule encoding pparticle of a preferred target region on a nucleic acid molecule encoding pparticle carrier. The antisense oligonucleotide comprises at least one modified nucleobase (which is a 5-methyl moiety) and at least one sugar moiety (preferably 2'-0-methoxyethyl moiety) and at least one sugar moiety (preferably 2'-0-methoxyethyl moiety) and at least one sugar moiety (preferably a 5-methyl cytosine). The antisense compounds are useful for treating cancer, osteoporosis, diabetes or various endocrine disorders. The Human ppart delta gene is located on chromosome 6p21. The present sequence is a human pparticle CDMA target
                                                                    Matches
                                                                                     Query Match
Best Local S
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                                                                                                                                            Sequence 20 BP; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antisense oligonucleotides of 8-80 nucleobases, useful for treating cancer, diabetes, osteoporosis or various endocrine disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to an antisense oligonucleotide comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 16; SEQ ID NO 218; 155pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-022078/02
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                                2073 GAGCCATCCAAAGAAACACT 2092
                                                                                       Similarity
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GAGCCATCCAAAGAAACACT 20
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                                                                    Conservative
                                                                                                                                                                             the antisense oligonucleotides of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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                                                                                       Score 20;
Pred. No.
                                                                    Mismatches
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                                                                    Indels
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RESULT 2 ADG86835/c

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ADG86835 standard; DNA; 20 BP

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TTCAGAGCAAAAGACTTGAG 2075

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The invention relates to an antisense oligonucleotide comprising 8-80 concleobases in length targeted to the coding region of a nucleic acid molecule encoding ppak-delta (peroxisome proliferative activated receptor delta), where the antisense compound inhibits the expression of the PPAR-delta (peroxisome proliferative activated receptor delta), where the antisense compound inhibits the expression of the PPAR-delta (peroxisome proliferative) and colds fully defined in the specification. Also included are a compound of 8-80 nucleobases in length that specifically hybridises with at least an 8-nucleobase portion of a preferred target region on a nucleic acid molecule encoding PPAR-delta and a composition comprising the antisense oligonucleotide and a carrier. The antisense oligonucleotide comprises at least one modified internucleoside linkage (preferably a phosphorothioate linkage), at least one sugar molety (preferably 2'-0-methoxyethyl molety) and at least one modified nucleobase (which is a 5-methyl cytosine). The antisense compounds are useful for treating cancer, osteoporosis, diabetes or various endocrine disorders. The Humann PPAR delta gene is located on chromosome 6p21. The present sequence is an antisense oligonucleotide of the invention targeting human PPAR delta.
Query Match
Best Local Similarity
Matches 20; Conserv
                                                                Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; SEQ ID NO 71; 155pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense oligonucleotides of 8-80 nucleobases, useful for treating cancer, diabetes, osteoporosis or various endocrine disorders.
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16. .20
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/mod_ba
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/mod_ba
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/mod_base= OTHER
/note= "Phosphorothioate linkages and all cytidines are
_methylcytidines"
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/note= "2'-methoxyethyl residue"
                                                                    5 C;
                                  40.0%;
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               Score 20; Pred. No.
                                                                    G; 8 T; 0 U; 0 Other;
   Mismatches
                 DB 1
                                  1;
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                                 Length 20
     Indels
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   Gaps
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ARESULT 3
ADG869818
ID ADG8
XX ADG8
XX ADG8
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RESULT 4
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Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of a preferred target region on a nucleic acid molecule encoding PPAR-delta and a composition comprising the antisense oligonucleotide and a carrier. The antisense oligonucleotide comprises at least one modified internucleoside linkage (preferably a phosphorothioate linkage), at least one sugar moiety (preferably 2'-O-methoxyethyl moiety) and at least one modified nucleobase (which is a 5-methyl cytosine). The antisense compounds are useful for treating cancer, osteoporosis, diabetes or various endocrine disorders. The Human PPAR delta gene is located on chromosome 6p21. The present sequence is a human PPAR delta cDNA target sequence for the antisense oligonucleotides of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to an antisense oligonucleotide comprising 8-80 nucleobases in length targeted to the coding region of a nucleic acid molecule encoding PPAR-delta (peroxisome proliferative activated receptor delta), where the antisense compound inhibits the expression of the PPAR-delta and has any of the 66 sequences of 20 amino acids fully defined in the specification. Also included are a compound of 8-80 nucleobases in length that specifically hybridises with at least an 8-nucleobase portion
11-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense oligonucleotides of 8-80 nucleobases, useful for treating cancer, diabetes, osteoporosis or various endocrine disorders.
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                                                                 ADG86983;
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                                                                                                                            ADG86983 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC
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20; Conserv
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(first entry)
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                                                                                                                            CDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; **
                                                                                                                                                                                                                                                                                                                                                                                                                                                            40.0%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 4.6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Watt AT;
                                                                                                                                20
                                                                                                                                ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 at least
                                                                                                                                                                                                                                                                                                                                                                                                                             0
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Human; ss; PPAR delta; peroxisome proliferative activated receptor delta; antisense gene therapy; cytostatic; osteopathic; antidiabetic; cancer; osteoporosis; diabetes; endocrine disorder.
                                                                                                                                                                    WPI; 2004-022078/02.
                                                                                                                                                                                 Gaarde W,
                                                                                                                                                                                                           31-MAY-2002; 2002US-00160807
                                                                                                                                                                                                                         31-MAY-2002; 2002US-00160807
                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                   US2003224514-A1
                                                                                                                                                                                                                                                                                                        Human PPAR antisense oligonucleotoide target sequence #45
                                                                                                                                                                                               (ISIS-) ISIS PHARM INC
                                                                                                                                                                                 Freier SM,
                                                                                                                                                                                 AT;
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The invention relates to an antisense oligonucleotide comprising 8-80 cnucleobases in length targeted to the coding region of a nucleic acid molecule encoding PPAR-delta (peroxisome proliferative activated receptor delta), where the antisense compound inhibits the expression of the PPAR-cc delta and has any of the 66 sequences of 20 amino acids fully defined in the specification. Also included are a compound of 8-80 nucleobases in clearly hybridises with at least an 8-nucleobase portion of a preferred target region on a nucleic acid molecule encoding PPAR-cc delta and a composition comprising the antisense oligonucleotide and a cc carrier. The antisense oligonucleotide comprises at least one modified internucleoside linkage (preferably a phosphorothioate linkage), at least one sugar moiety (preferably 2'-O-methoxyethyl moiety) and at least one compounds are useful for treating cancer, osteoporosis, diabetes or various endocrine disorders. The Human PPAR delta gene is located on component for the present sequence is a human PPAR delta CDNA target sequence for the antisense oligonucleotides of the invention.

Example 16; SEQ ID NO 219; 155pp; English

New antisense oligonucleotides of 8-80 nucleobases, useful for treating cancer, diabetes, osteoporosis or various endocrine disorders.

Sequence 20 BP; 6 A; 6 C; 4 G; 4 T; 0 U; 0 Other;

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5
                            Query Match
Best Local S
                      Matches
2086 AAACACTAAGCTCTCTGGGC 2105
                             Similarity
                     40.0%;
llarity 100.0%;
Conservative
                     0
                            Score 20;
Pred. No.
                      Mismatches
                            DB 1;
                      0
                                   Length 20
                      Indels
                      0
                      Gaps
                      0
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ADG86836/c
ID ADG868
XX
AC ADG868
XX
DT 11-MAR
XX
DT 11-MAR
XX
WHuman
XX
KW Human;
KW antise
KW osteop
XX
OS Homo s
                                                                                                                                                            片
                                                                                                                                 RESULT 5
                                                                               11-MAR-2004
                                                                                                                ADG86836 standard; DNA;
                                                                              (first entry)
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antisense gene therapy; cytostatic; osteopa osteoporosis; diabetes; endocrine disorder. Human; ss; PPAR delta; peroxisome proliferative activated receptor delta; antisense gene therapy; cytostatic; osteopathic; antidiabetic; cancer;

Human PPAR antisense oligonuclotide ISIS 136915

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RESULT
ADG8683
ID AD
XX
AC AD
XX
AC AD
XX
DT 11
XX
DE Hu
XX
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                                                                                                                                                                                                                                                                                                                                                           Ś
                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  delta and a composition comprising the antisense oligonucleotide and a carrier. The antisense oligonucleotide comprises at least one modified internucleoside linkage (preferably a phosphorothioate linkage), at least one modified nucleobase (which is a 5-methoxyethyl moiety) and at least one modified nucleobase (which is a 5-methyl cytosine). The antisense compounds are useful for treating canner, osteoporosis, diabetes or various endocrine disorders. The Human PPAR delta gene is located on chromosome 6p21. The present sequence is an antisense oligonucleotide of the invention targeting human PPAR delta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to an antisense oligonucleotide comprising 8-80 nucleobases in length targeted to the coding region of a nucleic acid molecule encoding PPAR-delta (peroxisome proliferative activated receptor delta), where the antisense compound inhibits the expression of the PPAR-delta and has any of the 66 sequences of 20 amino acids fully defined in the specification. Also included are a compound of 8-80 nucleobases in length that specifically hybridises with at least an 8-nucleobase portion of a preferred target region on a nucleic acid molecule encoding PPAR-delta and accompanies of a preferred target region on a nucleic acid molecule encoding PPAR-delta and accompanies of a preferred target region on a nucleic acid molecule encoding PPAR-delta accompanies of a preferred target region on a nucleic acid molecule encoding PPAR-delta accompanies of a preferred target region on a nucleic acid molecule encoding PPAR-delta accompanies of a preferred target region on a nucleic acid molecule encoding PPAR-delta acid molecule encoding PPAR-delta acid molecules acid molecule encoding PPAR-delta acid molecules acid
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modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New antisense oligonucleotides of 8-80 nucleobases, useful for treating cancer, diabetes, osteoporosis or various endocrine disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaarde W,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-MAY-2002; 2002US-00160807
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                                                                              11-MAR-2004
                                                                                                                               ADG86837;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20
                        Human PPAR antisense oligonuclotide ISIS 136916
                                                                                                                                                                                   ADG86837 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                           2073 GAGCCATCCAAAGAAACACT 2092
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Freier SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         B₽;
                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /mod_base= OTHER
/note= "Phosphorothioate linkages and all cytidines are
-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
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/note= "2'-methoxyethyl residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                note= "2'-methoxyethyl residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   *tag=
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                                                                                                                                                                                                                                                                                                                                                                                                       40.0%; Dre
100.0%; Pre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          . 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3 C; 6 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 20;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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RESULT 7
ADL34920
ID ADL3
XX

ADL34920 standard; DNA; 20

ВP

В Ś

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The invention relates to an antisense oligonucleotide comprising 8-80 nucleobases in length targeted to the coding region of a nucleic acid completule encoding PPAR-delta (peroxisome proliferative activated receptor delta), where the antisense compound inhibits the expression of the PPAR-celta and has any of the 66 sequences of 20 amino acids fully defined in the specification. Also included are a compound of 8-80 nucleobases in the specifically hybridises with at least an 8-nucleobase portion of a preferred target region on a nucleic acid molecule encoding PPAR-delta and a composition comprising the antisense oligonucleotide and a correct. The antisense oligonucleotide comprises at least one modified internucleoside linkage (preferably a phosphorothicate linkage), at least one sugar moiety (preferably 2'-o-methoxyethyl moiety) and at least one modified nucleobase (which is a 5-methyl cytosine). The antisense or compounds are useful for treating cancer, ostooporosis, diabetes or various endocrine disorders. The Human PPAR delta gene is located on compounds are present sequence is an antisense oligonucleotide of the invention targeting human PPAR delta.
                                                                                                        Query Match
Best Local :
                                                                                Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; ss; PPAR delta; peroxisome proliferative activated receptor delta; antisense gene therapy; cytostatic; osteopathic; antidiabetic; cancer; osteoporosis; diabetes; endocrine disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US2003224514-A1.
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                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; SEQ ID NO 73; 155pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense oligonucleotides of 8-80 nucleobases, useful for treating cancer, diabetes, osteoporosis or various endocrine disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaarde W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-MAY-2002; 2002US-00160807.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      31-MAY-2002; 2002US-00160807
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2086 AAACACTAAGCTCTCTGGGC 2105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sapiens.
                                                                                                        Similarity
                                                                                                                                                                                                 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Freier SM,
                                                                                                                                                                                              BP; 4 A; 4 C; 6 G; 6 T; 0 U; 0 Other;
                                                                            40.0%;
(larity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16. .20
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/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "Phosphorothicate linkages and all cytidines are 5
-methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         note= "2'-methoxyethyl residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           -methoxyethyl
                                                                                                        Score 20;
; Pred. No.
                                                                                o
;
                                                                                Mismatches
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                                                                                                           4.6;
                                                                                                                                          DB 1;
                                                                                0,
                                                                                                                                          Length 20;
                                                                                Indels
                                                                                ٠<u>,</u>
                                                                                Gaps
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WWW.SXEEEEEEEEEEEEEEEEKKEXEXEXEXEXEXEXEXSOOOOOOO

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RESULT 8
ADL34774/c
ID ADL347
XX
AC ADL347
XX
DT 17-JUN
XX
DE Antise
XX
W antise
XW antise
XW hyperp
XW primer
XX
XX primer
XX
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                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 20; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention describes novel antisense oligonucleotides targeted to a nucleic acid encoding PPAR-delta, which specifically hybridise to and inhibit expression of PPAR-delta. The oligonucleotide specifically hybridises with at least an 8-nucleobase portion of an active site on the nucleic acid molecule encoding the PPAR-delta and comprises at least one modified internucleoside linkage, which is a phosphorothioate linkage, at least one modified sugar moiety, which is a 2'-O-methoxyethyl sugar moiety or at least one modified nucleobase, which is a 5-methylcytosine. The antisense oligonucleotides are useful for preparing a composition for treating hyperproliferative disorders, e.g., cancer. The oligonucleotides of the inventor have disorders and can be used for cane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New antisense oligonucleotide, having a sequence targeted to acid encoding PPAR-delta, useful for preparing a composition hyperproliferative disorder, e.g., cancer.
                              antisense; PPAR-delta; human; hybridisation; inhibitor;
phosphorothioate linkage; 2'-O-methoxyethyl sugar; 5-methylcytosine;
hyperproliferative disorder; cancer; cytostatic; gene therapy; ss;
                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 16; SEQ ID NO 218; Opp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GAAR/) GAARDE W.
(FREI/) FREIER S M.
(WATT/) WATT A T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADL34920;
                                                                                                 Antisense
                                                                                                                                                                                                ADL34774 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               the invention have cytostatic activity and can be used for gene
                                                                                                                                                                                                                                                                                                              2073 GAGCCATCCAAAGAAACACT 2092
                                                                                                                                                                                                                                                                               GAGCCATCCAAAGAAACACT 20
                                                                                                   oligonucleotide ISIS 136915
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Freier SM,
                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                             A; 6 C; 3 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                 40.0%; Sec. No. 100.0%; Pred. No. 100.0%; Mismatches
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                                                                                                                                                                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                             Length 20
                                                                                                                                                                                                                                                                                                                                              Indels
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                                                                                                                                                                                                                                                                                                                                                Gaps
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RESULT 9
ADD3477/c
ID ADD347
XX
AC ADD347
AC ADD347
XX
DT 17-JUN
XX
DE Antise
XX
Antise
Antise
XX
Antise
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                  antisense; PPAR-delta; human; hybridisation; inhibitor;
phosphorothioate linkage; 2'-O-methoxyethyl sugar; 5-methylcytosine;
hyperproliferative disorder; cancer; cytostatic; gene therapy; ss;
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                               31-MAY-2002; 2002US-00160807
                                                                                           05-SEP-2003; 2003US-00655847
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Best Local S
Matches 20
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New antisense oligonucleotide, having a sequence targeted to acid encoding PPAR-delta, useful for preparing a composition hyperproliferative disorder, e.g., cancer.
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                                                                                                                                                                                                                                                                                                     antisense; PPAR-delta; human; hybridisation; inhibitor;
phosphorothioate linkage; 2'-O-methoxyethyl sugar; 5-methylcytosine;
hyperproliferative disorder; cancer; cytostatic; gene therapy; ds.
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                                                       WPI; 2004-282460/26.
                                                                                                                                                                   31-MAY-2002; 2002US-00160807
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Pred. No.
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Best Local :
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                                                                                                                                                                                                                                                                                                                            New antisense oligonucleotide, having a sequence acid encoding PPAR-delta, useful for preparing a
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phosphorothioate linkage; 2'-O-methoxyethyl sugar; 5-methylcytosine;
hyperproliferative disorder; cancer; cytostatic; gene therapy; ds.
                                                                                                                                                                                                                                                                                                hyperproliferative disorder, e.g.,
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Pred. No.
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nucleic acid encoding PPAR-delta, which specifically hybridise to and inhibit expression of PPAR-delta. The oligonucleotide specifically hybridises with at least an 8-nucleobase portion of an active site on the nucleic acid molecule encoding the PPAR-delta and comprises at least one modified internucleoside linkage, which is a phosphorothioate linkage, at least one modified sugar moiety, which is a 2'-O-methoxyethyl sugar moiety or at least one modified nucleobase, which is a 5-methoxyethyl sugar

Example 16;

SEQ ID NO 217;

Opp; English.

invention

describes novel

antisense oligonucleotides targeted to

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RESULT 12
ADL34775/c
ID ADL347
Query Match
Best Local Similarity 100.
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phosphorothioate linkage; 2'-O-methoxyethyl sugar; 5-methylcytosine;
hyperproliferative disorder; cancer; cytostatic; gene therapy; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADL34775
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADL34775 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The antisense oligonuclectides are useful for preparing a composition for treating hyperproliferative disorders, e.g., cancer. The oligonuclectides of the invention have cytostatic activity and can be used for gene
                                                                                                                  Sequence 20 BP; 4 A; 4 C; 6
                                                                                                                                                                                                                                                                                                                                                                                                 This invention describes novel antisense oligonucleotides targeted
                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 15; SEQ ID NO 73; Opp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GAAR/) GAARDE W. (FREI/) FREIER S M. (WATT/) WATT A T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-SEP-2003; 2003US-00655847
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  hyperproliferative disorder, e.g., cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-282460/26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2056 TTCAGAGCAAAAGACTTGAG 2075
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                      40.0%; br
100.0%; Pr
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100.0%; Pri
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                                                                                                                  G; 6 T; 0
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                                                   Score 20;
Pred. No.
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Pred. No.
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                                                                                                                  U; 0 Other;
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Query Match Best Local Similarity

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                                                          which distinguishes between methylated and non-methylated Cpg
dinucleotides within the target nucleic acid. ABZ09861 to ABZ1118
represent specifically claimed nucleotide sequences from the present
invention. Oligonucleotides from the present invention can be used: for
differentiating between healthy haematopoietic cells and proliferative
disorder haematopoietic cells; for differentiating between acute
lymphocytic leukaemia and acute myelogenous leukaemia; as probes for
determining the cycosine methylation state and/or single nucleotide
polymorphisms (SNPs) of haematopoietic cell proliferation disorder
related sequences and their complements; and as primers for the
amplification of haematopoietic cell proliferation disorder related sequences. The nucleotide sequences from the present invention can also
sequences. The nucleotide sequences from the present invention can also
be used for detecting a predisposition to, differentiation between
subclasses, diagnosis, prognosis, treatment and/or monitoring of
haematopoietic cell proliferative disorders. The present method enables a
highly specific classification of haematopoietic cell proliferative
disorders allowing for improved and informed treatment of patients
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; ; ; <sub>20</sub>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes a method for detecting and differentiating between haematopoietic cell proliferative disorders associated with at least 1 gene and/or their regulatory regions in a subject. The method comprises contacting a target nucleic acid in a biological sample obtained from the subject with at least 1 reagent,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Detecting and differentiating between hematopoietic cell proliferative disorders, comprises contacting a target nucleic acid with a reagent that distinguishes between methylated and non-methylated CpG dinucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Berlin K, Braun A, Distler J, Guetig D, Olek A, Piepenbrock C, Adorjan P, Grabs Lewin A, Lipscher E, Maier S, Model F, Schwope I, Ziebarth H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; haematopoietic cell proliferation disorder; cytostatic; gene therapy; lymphocytic leukaemia; acute myelogenous leukaem cytosine methylation state; probe; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 15; Page 59; 117pp; English
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Mueller V, Otto
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                                                                                                                                                        The invention relates to a novel method for detecting and differentiating between colon cell proliferative disorders associated with at least one gene or its regulatory regions. The method comprises contacting a target nucleic acid in a biological sample obtained from the subject with at least one reagent or a series of reagents, where the reagent or series of reagents, where the reagent or series of dinucleotides within the target nucleic acid. The molecules of the invention demonstrate cytostatic activity whilst the method may useful for detecting and differentiating between colon cell proliferative disorders, including cancers such as colon adenoma and colon carcinoma. The PNA (peptide nucleic acid) -oligomers are useful as probes for determining cytosine methylation state or single nucleotide polymorphisms. The current sequence is that of the hybridisation of the invention which was used to analyse the genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Adorjan E
Rujan T,
                                                                                                                 Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                            Detecting and differentiating between colon cell proliferative disorders associated with a gene or its regulatory regions comprises contacting a target nucleic acid in a biological sample obtained from the subject with
                                                                                                                                                                                                                                                                                                                                                                                  Claim 36; Page 42; 74pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            colon cell proliferative disorder; non methylated CpG dinucleotide; cytostatic; cancer; adenoma; carcinoma; cytosine methylation state;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-731620/69.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27-FEB-2002; 2002EP-00004551.
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Human lymphoid cell proliferative disorder gene CpG analysis oligo #218.

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ID ADC70291 standard; DNA; 11

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AC ADC70291;

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DT 18-DEC-2003 (first entry

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PCR; primer; ss; lung cel

XW

PCR; primer; ss; lung cel

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Adenocarcinoma; squamous

XW

Cytosine methylation stat

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W02003052135-A2.

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(EPIG-) EPIGENOMICS AG.

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PI Burger M, Field JK, Gen

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PI Burger M, Field JK, Gen

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PI Burger M, Field JK, Gen

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PT Useful for diagnosing, tr

Cell proliferative disord

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Claim 15; SEQ ID NO 781;

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CC This invention relates to differentiating between me cc differentiate and diagnos of the least one gene and/or refers to a method comprise to biological sample with at the diagnosing colloquidal problemanum colloq
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                                                         29-JAN-2004
                                                                                                                  ADE84512;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          differentiating between lung cell proliferative disorders associated with at least one gene and/or their regulatory regions. Specifically, it refers to a method comprising contacting a target nucleic acid in a biological sample with at least one reagent, wherein the reagent is able to distinguish between methylated and non-methylated CpG dinucleotides present in the target DNA. As such, it is possible to further differentiate and diagnose medical conditions including adenocarcinoma and squamous cell carcinoma, and their respective adjacent lung tissue. The present invention describes cytostatic oligomers and PNA-oligomers that are useful as probes for determining the cytosine methylation state or single nucleotide polymorphisms (SNPs) of the target sequence. This oligonucleotide sequence is a primer oligomer used for the analysis of CpG positions within genomic DNA, used in an exemplification of the
                                                                                                                                                                                 ADE84512 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Detecting and differentiating cytosine methylation state of genomic DNA, useful for diagnosing, treating prognosticating and/or monitoring lung cell proliferative disorders e.g. adenocarcinoma and squamous cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR; primer; ss; lung cell proliferative disorder; CpG dinucleotide; adenocarcinoma; squamous cell carcinoma; cytostatic; probe; PNA-oligomer; cytosine methylation state.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention relates to a novel method for detecting and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-DEC-2001; 2001DE-01061625
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                                                                                                                                                                                                                                                                                                                                                                                         2079 TCCAAAGAAACACTAA 2094
                                                                                                                                                                                                                                                                                                                                   16 TCCAAACAAACACTAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 3 A; 0 C; 5 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                  (first entry)
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                                                                                                                                                                                 DNA; 18
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Pred. No. 20;
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RESULT 17
AAX75280
ID AAX75
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AC AAX75
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DT 28-JU
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                                                                                                                                                                                                                                                                                                                                                                    CC between lymphoid cell proliferative disorders associated with at least CC one gene and/or their regulatory regions in a subject by contacting a CC target nucleic acid in a biological sample obtained from the subject with CC at least one reagent or series of reagents that distinguish between CC acid. The genes and/or their regulatory regions are preferably selected from MDR1, CSNK2B, EGR4, AR, CDK4, RB2, CDC32A, GPID beta, MVOD1, CDH3, CC GSTP1, HIC-1, MGMT, MLH1, MOS, MYC, PTEN, RBL2, TGFBR2, TCH3, CC GSTP1, HIC-1, MGMT, MLH1, MOS, MYC, PTEN, RBL2, TGFBR2, CDKN2B, FOS, CC GSK1beta, ESR1, APAF1, BAK1, BAX or HOXA5. Oligomers, peptide nucleic acid (PNA)-oligomers and/or isolated nucleic acids based on the sequences of the genes are useful for detecting the methylation state of all the CC Godinucleotides within one or more the sequences, or their complements, conditions such as diffuse large B-cell lymphoma, mantle cell lymphoma, cuch as diffuse large B-cell lymphoma and follicular conditions such as diffuse large B-cell lymphoma and follicular complements an oligonucleotide cell proliferative disorder. This sequence created and or supplements and consistion between subclasses, diagnosis, treating and/or above mention between subclasses, diagnosis, prognosis, treating and/or above mention between subclasses, diagnosis, prognosis, treating and/or above mention between subclasses diagnosis prognosis treating and/or above mentions.
                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Detecting and differentiating between lymphoid cell proliferative disorders comprises contacting a target nucleic acid with at least one reagent that distinguishes between methylated and non-methylated CpG dinucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               lymphoid cell proliferative disorder; methylation; methylated CpG dinuclectide; single nuclectide polymorphism; diffuse large B-cell lymphoma; mantle cell lymphoma; chronic lymphocytic leukemia; small lymphocytic lymphoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a method of detecting and differentiating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 30; SEQ ID NO 508; 448pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-457621/43
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28-DEC-2001;
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28-JUL-1999
                                     AAX75280;
                                                                        AAX75280 standard;
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                                                                                                                                                                                           2079 TCCAAAGAAACACTAA 2094
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                                                                                                                                                                                                                                                              Similarity
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                                                                                                                                                                                                                                              Conservative
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2001DE-01064501.
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                                                                                                                                                                                                                                                                                                                     G; 10 T; 0 U; 0 Other;
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Pred. No. 2
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RESULT 18
AAF47731/c
ID AAF477
XX AAF477
XX AAF477
XX 30-MAR
XX IGFBP3
XX ANTISE
KW ANTISE
KW Skin c
KW IGF bj
KW GYOWL
KW GYOWL
KW GYOWL
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Best Local S
Matches 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention
               Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatological; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleic acid molecule modulating VEGF receptor(s) gene expression or stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17
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                                                                                                                                                   IGFBP3 oligonucleotide #1151.
                                                                                                                                                                                   30-MAR-2001
                                                                                                                                                                                                                                                   AAF47731 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 4; Page 179; 218pp; English.
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11-JAN-1996;
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96US-00584040.
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Pred. No. 22;
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RESULT 19
AAF47730/c
ID AAF477
XX AAF477
XX AAF477
XX 30-MAR
XX IGFBP3
XX Antise
KW Cytost
KW Skin d
KW IGF Moderato
KW IGF Moderato
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Best Local (
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                                                               Antisense therapy; antiproliferative; antiinflammatory; antipsoriatic; cytostatic; dermatclogical; cardiant; virucide; ophthalmological; keloid; skin disorder; Insulin-like Growth Factor I receptor; IGF-1; pityriasis; IGF binding protein; IGFBP-2; IGFBP3; inflammation; psoriasis; pilaris; growth factor mediated cell proliferation; ichthyosis; serborrhoea; ruba; keratosis; neoplasia; scleroderma; wart; skin cancer; sclerotic disease; hyperneovascular condition; hyperplasia; kidney disease; hyperacovascular condition of the retina; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to a method for ameliorating the effects of skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF] 1 receptor, IGF binding protein [IGFBP] 2 or IGFBP3), which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as neovascular condition of the retina, warts the factor of the skin, a hyperneovascular condition such as neovascular condition of the retina,
                                                                                                                                                                                                                                                                   30-MAR-2001
                                                                                                                                                                                                                                                                                                       AAF47730;
                                                                                                                                                                                                                                                                                                                                        AAF47730 standard; DNA; 15 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ameliorating the effects of a disorder, e.g. psoriasis, by administer UV (ultra-violet) treatment (optional) and an antisense nucleic acid inhibits or reduces growth factor mediated cell proliferation and/or
                                                                                                                                                                                                                                  IGFBP3 oligonucleotide #1150.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               2076 CCATCCAAAGAAAC 2089
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92.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Edmondson
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 12.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Η,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    by administering nucleic acid that
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WO200078341-A1

WO9523225-A2

Mus musculus.

domo sapiens

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RESULT 20
AAT52513/c
ID AAT525
XX AAT525
XX AC AAT525
XX 10-APR
DT 10-APR
XX 10-APR
XX MOUSE
XW Enzyma
KW Enzyma
KW Interc
KW Interc
KW TW-all
KW Transl
KW Philad
KW Philad
KW Philad
XX myocar
KW myocar
KW myocar
KW ms.
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Best Local S
Matches 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           skin disorders. The method comprises contacting the skin with an antisense oligonucleotide, (for Insulin-like Growth Factor [IGF]-1 receptor. IGF binding protein [IGFBP]-2 or IGFBP], which is capable of inhibiting or reducing growth factor mediated cell proliferation, inflammation and/or other disorders. The present sequence is an oligonucleotide which can be used to design the antisense oligonucleotides of the present invention (see AAF45151 and AAF45153-F45161). The method is useful for ameliorating the effects of psoriasis, ichthyosis, pityriasis, ruba, pilaris, serborrhoea, keloids, keratosis, neoplasias, scleroderma, warts, benign growths, cancers of the skin, a hyperneovascular condition such as a neovascular condition of the retina, brain or skin, growth factor-mediated malignancies, other sclerotic disease, kidney disease, hyperproliferation of the inside of blood vessels or any other hyperplasia
                                                                                                                                                                                           Enzymatic nucleic acid; ribozyme; trans cleavage; inhibition; gene expression; downregulation; interleukin-5; IL-5; ICAM-1; intercellular adhesion molecule; rel A; tumour necrosis factor; TNF-alpha; respiratory syncytial virus; RSV; bcr-abl; oncogene; translocation; chronic myelogenous leukaemia; CML; cancer; translocation; chronic myelogenous leukaemia; CML; cancer;
                                                                                                                    atherosclerosis; myocardial infarction; stroke; restenosis; transplant rejection; rheumatoid arthritis; psoriasis; myocardial ischaemia; Kawasaki disease; septic shock; HIV;
                                                                                                                                                                             Philadelphia chromosome; inflammation; autoimmune disease;
                                                                                                                                                                                                                                                                                                                 Mouse ICAM hammerhead ribozyme target sequence (nt. position 2593).
                                                                                                                                                                                                                                                                                                                                                        25-MAR-2003
10-APR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                 AAT52513;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT52513 standard; RNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to a method for ameliorating the effects
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 7; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ameliorating the effects of a disorder, e.g. psoriasis, by administering UV (ultra-violet) treatment (optional) and an antisense nucleic acid that inhibits or reduces growth factor mediated cell proliferation and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-041421/05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MURD-) MURDOCH CHILDRENS RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2076 CCATCCAAAGAAAC 2089
                                                                                                 immunodeficiency virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CCATTCAAAGAAAC 2
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                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                          revised)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP
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Pred. No. 27;
                                                                                                 acquired
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1; Length 15;
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                                                                                                 deficiency syndrome; AIDS;
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31-AUG-1995

95WO-IB000156.

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                                            Query Match
Best Local Similarity
Matches 12; Conserv
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03-OCT-1994;
07-OCT-1994;
11-OCT-1994;
04-NOV-1994;
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Modak A,
Tracz D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-AUG-1994;
19-AUG-1994;
19-AUG-1994;
02-SEP-1994;
08-SEP-1994;
23-SEP-1994;
23-SEP-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-FEB-1994;
29-MAR-1994;
04-APR-1994;
07-APR-1994;
15-APR-1994;
                                                                                                                  The present sequence represents a preferred target sequence for an enzymatic nucleic acid (i.e. a ribozyme) which cleaves ICAM-1 mRNA at the nucleotide base position indicated in the DE line. Regions of the mRNA that do not form secondary folding structures and that contain potential hammerhead and hairpin ribozyme cleavege sites were identified by computer analysis. Ribozymes directed against these mRNA sequences were designed and synthesised with modifications that improve their nuclease resistance. The ribozymes cleave the ICAM-1 target sequences and thereby inhibit ICAM-1 expression, making them useful for reducing transplant rejection and alleviating symptoms in patients with rheumatoid arthritis, asthma and other inflammatory disorders. (Updated on 25-MAR-2003 to
                                                                                              Sequence 15
                                                                                                                                                                                                                                                                                   Claim
                                                                                                                                                                                                                                                                                                          Ribozymes having modified bases and methods for producing them in inhibiting disease related genes.
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                                                                                                                                                                                                                                                                                                                                                                                                           Stinchcomb
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28-NOV-1994
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18-MAY-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                  (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-AUG-1994
                       2058 CAGAGCAAAAGA 2069
14
                                                                                                                                                                                                                                                                                  2; Page 180; 407pp; English
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Pavco P,
Usman N,
                                                                                                                                                                                                                                                                                                                                                                                                             Ğ.
                                                                                             BP; 0
                                               Conservative
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94US-00291932.
94US-00291
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94US-00319492.
94US-00321993.
94US-00331847.
94US-00337608.
94US-00345516.
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94US-00293520
94US-0030000
94US-0030030
94US-00311486
94US-00311749
94US-00314397
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94US-00363233.
95US-00380734.
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94US-00271280.
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                                                                                             A;
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Beigleman L,
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                                            24.0%; Score 12; DB 1; Length 15; 100.0%; Pred. No. 31; tive 0; Mismatches 0; Indels
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                                                                                             C; 2 G; 0 T; 8 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                  nzo A, Draper KG, Dudycz LW;
Matulic-Adamic J, Mcswiggen JA;
Sullivan SM, Sweedler D, Thompson JD;
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                                              Gaps
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RESULT 21 ABF44787 ID ABF44

ABF44787 standard; DNA; 13

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RESULT 22
ABC05028/c
ID ABC050
XX
AC ABC050
XZ
DT 20-FEB
XX
Oligon
XX
SNP; 8
KW SNP; 8
KW peptid
KW centra
XX
OS Homo 8
XX
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Best Local S
Matches 12
                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; peptide nucleic acid; cytosine methylation; cardiovascular; primer; central nervous system; gastrointestinal; respiratory; immune; metal
                                                                                                                                                                                                                                                                                                                                                                                                                                               range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NoTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Set of oligonucleotides, useful for diagnosis and cell typing; designed to detect single-nucleotide polymorphisms and cytosin methylation status.
                    Homo sapiens
                                                                                                           Oligonucleotide SEQ ID NO 5019 for detecting SNP TSC0001741.
                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 13 BP; 7 A; 5 C; 0 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; peptide nucleic acid; cytosine methylation; central nervous system; gastrointestinal; re
                                                                                                                                          20-FEB-2002
                                                                                                                                                                      ABC05028
                                                                                                                                                                                                    ABC05028 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; SEQ ID NO 144784; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                               ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-OCT-2001.
                                                                                                                                                                                                                                                                                                            2077 CATCCAAAGAAAC 2089
                                                                                                                                                                                                                                                                                                                                         12;
                                                                                                                                                                                                                                                                                                                                                        Similarity
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                                                                                                                                                                                                                                                                              CATCCAAACAAAC 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Piepenbrock C,
                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                                          (first
                                                                                                                                                                                                    DNA;
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                                                                                                                                                                                                                                                                                                                                                       Score 11.4;
Pred. No. 30;
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                                                                                                                                                                                                                                                                                                                                                                     Length 13;
                                                                                                                                                                                                                                                                                                                                          Indels
                                                PNA; cancer; CNS; ular; primer; ss; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cytosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ilar; primer; ss;
immune; metabolic.
                                                                                                                                                                                                                                                                                                                                         0,
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RESULT 23
ABC05029
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Olek A,
WPI; 2001-657177/75
                       Olek A,
                                                                          07-APR-2000; 2000DE-01019173.
                                                                                                 06-APR-2001; 2001WO-IB000713
                                                                                                                           18-OCT-2001.
                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                   central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                                                                                                                                                 peptide
                                                                                                                                                                                                                                                     Oligonucleotide
                                                                                                                                                                                                                                                                                                       ABC05029;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       designed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Set of oligonucleotides, useful for diagnosis and cell typing, is
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                                                                                                                                                                                                              single nucleotide polymorphism; human; diagnosis; PNA; ide nucleic acid; cytosine methylation; cardiovascular;
                                                                                                                                                                                                                                                                                                                                                                                                                  2082 AAAGAAACACTAA 2094
                                                                                                                                                                                                                                                                                                                                                                                            13
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12; Conserv
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                       Piepenbrock C,
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                                                                                                                                                                                                                                                                             (first entry)
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                                                                                                                                                                                                                                                  SEQ ID NO 5020 for detecting SNP TSC0001741.
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Matches 12
                                                                                                                                                                          range of diseases including immune system, gastrointestinal, respirate central nervous system, cardiovascular and metabolic disorders. The coligomers are also used for detecting cell type differentiation. ABCO-ABCO9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI0010-ABH802073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                             acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNI and cytosine methylation status in chemically pretreated genomic DNA. I oligonucleotides are used for diagnosis and/or prognosis of cancer and
                                                                                                                                                                                                                                                                                                                                                                                                          Set of oligonucleotides, useful for diagnosis and designed to detect single-nucleotide polymorphisms methylation status.
                                                                                                                             Sequence 13 BP; 9 A; 3 C; 0 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                              Claim
                                                                                                                                                          ftp.wipo.int/pub/published_pct_sequences
                             2082 AAAGAAACACTAA 2094
                                                                                                                                                                                                                                                                                                                                             invention
                                                             12;
                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                SEQ ID
                                                               Conservative
                                                                                                                                                                                                                                                                                                                                             describes novel oligonucleotide primers or peptide nucleic
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                                                                             Score 11.4;
Pred. No. 30;
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ABF48836/c
ID ABF488
XX ABF4888
XX ABF488
XX ABF488
XX Oligon
XX SNP; s
KW Peptid
KW Centra
XX WO2001
XX WO2001
XX WO2001
XX WO2001
XX Of-APF
XX O SNP; single nucleotide polymorphism; human; peptide nucleic acid; cytosine methylation; Homo sapiens. Oligonucleotide 21-FEB-2002 ABF48836 ABF48836 standard; DNA; 13 nervous (first SEQ system; gastrointestinal; ä entry) NO 148833 for ВP detecting diagnosis; PNA; cancer; CNS; cardiovascular; primer; ss; espiratory; immune; metabolic SNP TSC0037567

WO200177384-A2.

06-APR-2001; 2001WO-IB000713. 18-OCT-2001

07-APR-2000; 2000DE-01019173.

(EPIG-) EPIGENOMICS ÃĢ.

Piepenbrock C, Berlin ۲

WPI; 2001-657177/75.

methylation status. Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine

Claim 1; SEQ ID NO 148833; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, cardiovascular

oligomers are also used for detecting cell type differentiation. ABC00010 -ABC99989, ABF00010-ABF99989, ABH00010-ABF99989, and ABF00010-ABF32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at

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ABF44786; babr44786; plb 48F44786; plb 48F42001; plb 48F4201-657177/75. plb 48F4201-657177/
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                                                                                                                                                                                                                                                                                                                                                                                                         acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; SEQ ID NO 144783; 29pp + Sequence Listing; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosin
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                                                                                                                   Sequence 13 BP; 1 A; 0 C; 5
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ide nucleic acid; cytosine methylation; cardiovascular;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     invention describes novel oligonucleotide primers or peptide nucleic
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Pred. No. 30;
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ABF48837
ID ABF48
XX ABF48
AC 
RESULT 27
ABF26976/c
ID ABF269
XX
AC ABF269
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AC ABF269
XX
DT 21-FEB
XX
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                                   21-FEB-2002
                                                                                                                                                      ABF26976 standard; DNA; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 13 BP; 2 A; 6 C; 1 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-APR-2000; 2000DE-01019173.
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Pred. No. 30;
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RESULT 28
ABF26977
ID ABF26
XX ABF26
XX ABF26
XX Oligo
DT 21-FE
XX Oligo
XX SNP;
KW SNP;
KW Septi
KW Centr
XX Centr
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Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.

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               This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic form the invention.
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RESULT 31
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                                                                                                                                                                                                                                                                                        This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH99989, appresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; SEQ ID NO 185992; 29pp + Sequence Listing; German
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1 RAACAAACACTAA 13
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acid; cytosine methylation; cardiovascular; primer; ss;
system; gastrointestinal; respiratory; immune; metabolic
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Pred. No.
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Pred. No.
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RESULT 33
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ABF87457
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; peptide nucleic acid; cytosine methylation; cardiovascular; central nervous system; gastrointestinal; respiratory; immur
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                           Oligonucleotide
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                                                                                                                                                                                                                                                                                                                   ABI58987 standard; DNA; 12
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                                                                                                                       primer SEQ ID NO 358960 for detecting SNP TSC0009167.
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      immune; metabolic
                             cancer; CNS;
primer; ss;
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SNP; is
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Best Local Similarity
Matches 11; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                              07-APR-2000; 2000DE-01019173.
                                                                          06-APR-2001; 2001WO-IB000713.
                                                                                                          18-OCT-2001
                                                                                                                                       WO200177384-A2
                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                            Oligonucleotide primer SEQ ID NO 275965 for detecting SNP TSC0004052.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                AAAAAACACTAA 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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Pred. No. 35;
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(EPIG-) EPIGENOMICS

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Olek A,

Piepenbrock C,

Berlin K;

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RESULT 35
AB144086
ID AB144
XX AB144
XX Oligo
XX SNP;
XW Pepti
XW Centr
XX Centr
XX Homo
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; peptide nucleic acid; cytosine methylation; cardiovascular; central nervous system; gastrointestinal; respiratory; immur
                                                                                                                                                                                                                                                                                                                                                   Olek A,
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                                                                                                                      Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-APR-2000; 2000DE-01019173
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                                                                                                             ID NO 344059; 29pp + Sequence Listing; German
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                                                                                                                                                                                                                                                                                                                                                         Berlin K;
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                                                                                                                                                                                                       cytosine
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primer; 88;
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The

DNA.

Query Match

Sequence 12 BP; 3 A; 3 C; 1 G; 5 T; 0 U; 0 Other;

Score 10.4;

DB 1; Length 12;

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RESULT 36
ABI74566/c
ID ABI745
XX ABI745
XX ABI745
XX Oligon
XX SNP; s
KW Peptid
XX Centra
XX SNP; s
KW Centra
XX Homo s
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Best Local Simi
Matches 11;
                                                                                               This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolli disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 ABC9989, ABF00010-ABF9989, ABF00010-ABF9989, ABH00010-ABF9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; SEQ ID NO 374539; 29pp + Sequence Listing; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    methylation status
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                                                                      ftp.wipo.int/pub/published_pct_sequences
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Pred. No. 3
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RESULT 37
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Best Local Similarity
                                                                                                                                                                               This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                Sequence 12 BP; 7
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ABI40361;
                     ABI40361 standard; DNA; 12
                                                                                                                                                                                                                                                                                                                                Claim 1; SEQ ID NO 372985; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-657177/75
                                                                                                                                                                                                                                                                                                                                                                                                                                            (EPIG-)
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                                                                                               2078 ATCCAAAGAAAC 2089
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                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                A; 3 C; 0 G;
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                     BP.
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                                                                                                                     0; Mismatches
                                                                                                                               Score 10.4;
Pred. No. 35;
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0; Mismatches
                                                                                                                                                                2 T; 0 U; 0 Other;
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                                                                                                                                          DB 1;
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                                                                                                                                          Length 12;
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WO200177384-A2

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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomers are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oligonucleotide primer SEQ ID NO 340334 for detecting SNP
                                                                                                              SNP; single nucleotide polymorphism; human; peptide nucleic acid; cytosine methylation; central nervous system; gastrointestinal; re
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 12 BP; 7 A; 4 C; 0 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Olek A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-APR-2000; 2000DE-01019173.
                                                                                                                                                                                                           Oligonucleotide primer SEQ ID
                                                                                                                                                                                                                                                           22-FEB-2002
                                                                                                                                                                                                                                                                                                          ABI54654;
                                                                                                                                                                                                                                                                                                                                                        ABI54654 standard;
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                                                                      Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CAAACAAACACT
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                                                                                                                                                                                                                                                           (first
                                                                                                                                                                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                           entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 10.4;
Pred. No. 39
                                                                                                                                                                                                                  NO 354627 for detecting SNP TSC0049190.
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                                                                                                                     n; diagnosis; PNA; cancer; CNS; n; cardiovascular; primer; ss; respiratory; immune; metabolic
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cytosine
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RESULT 40
ABI61429
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Best Local Similarity
Matches 11; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                range of diseases including immune system, gastrointestinal, respirator central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC000-ABC99989, ABF00010-ABE99989, ABH00010-ABH9998 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, buwas obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
                                                       Olek A,
                                                                                                                                                                                               WO200177384-A2
                                                                                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                                                                                                                                                                                                                                       Oligonucleotide primer SEQ ID NO 361402 for detecting SNP TSC0052625
                                                                                                                                                                                                                                                                                                                                     22-FEB-2002
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Set of oligonucleotides,
                            WPI; 2001-657177/75
                                                                                                            07-APR-2000; 2000DE-01019173.
                                                                                                                                      06-APR-2001; 2001WO-IB000713.
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                                                                                  (EPIG-)
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                                                                                  EPIGENOMICS AG
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                                                       Piepenbrock C,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20.8%;
useful for diagnosis and cell typing,
                                                       Berlin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          G; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pred. No. 35;
); Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 10.4;
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RESULT 41
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           range of diseases including immune system, gastrointestinal, respirator central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC000-ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, buwas obtained in electronic format from WIPO at
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          range of diseases including immune system, gastrointestinal, recentral nervous system, cardiovascular and metabolic disorders oligomers are also used for detecting cell type differentiation
                                                             This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                              Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                  Claim 1;
                                                                                                                                                                                                                                                    WPI; 2001-657177/75.
                                                                                                                                                                                                                                                                                                                                                   07-APR-2000; 2000DE-01019173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oligonucleotide primer SEQ ID NO 328912 for detecting SNP TSC0034651.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABI28939
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 12 BP; 8 A; 3 C; 0 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                SEQ ID NO 328912; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                 Piepenbrock C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         standard;
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Pred. No. 35
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                                               respiratory,
             ABC00010
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-ABC99989, ABF00010-ABF99989,

ABH00010-ABH99989 and

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ABIGGG35/c
ID ABIGGG
XX ABIGGG
XX ABIGGG
XX ABIGGG
XX Oligon
XX SNP; kg
KW Peptid
XX VOIGON
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XX OIGON
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                                                                                    Query Match
Best Local S
Matches 11
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Matches 11
                                                                                                                                                                                                                                                                             acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cyrosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for disgnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                               Sequence 12 BP; 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 368608; 29pp + Sequence Listing; German.
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Pred. No. 35
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Oligonucleotide primer SEQ

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RESULT 45
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                                                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; peptide nucleic acid; cytosine methylation; cardiovascular;
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 07-APR-2000; 2000DE-01019173.
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peptide nucleic acid; cytosine methylation; cardiovascular;
                                                                      18-OCT-2001
                                                                                                        WO200177384-A2
                                                                                                                                                                                                                                                                                                                                               ABH88737 standard;
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                                                                                                                                                                        system;
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RESULT 46
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                                                 Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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Claim 1; SEQ ID NO 378848; 29pp + Sequence Listing; German

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RESULT 47
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range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                        This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
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                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
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                                                                                                                                                                              Oligonucleotide primer SEQ ID NO 327968 for detecting
                                                                                                                                                                                                                                             22-FEB-2002
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                                                                                                                                                   07-APR-2000; 2000DE-01019173
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Olek A,

Piepenbrock C,

Berlin

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosts and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory,
                                                                                                                                      Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              designed to detect methylation status.
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                                                                                                    Claim 1; SEQ ID NO 320231; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                  This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a oligonucleotides are used for diagnosis and/or prognosis of cancer and a oligonucleotides.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
Sequence 12
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Pred. No. 35;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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22-FEB-2002
                                                                                                                         ABI65055 standard; DNA; 12 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 12 BP; 1 A; 0 C; 3
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                                                            ABI65055;
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Pred. No. 35;
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                     H-ras; wild-type; immobilising; diagnosis; ethylene acrylic acid; ethylene methacrylic acid; polypropylene; biotin; cystic fibrosis
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                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                               One from an array of 58 cystic fibrosis oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                          01-JUL-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV06890
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                designed to detect methylation status.
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Best Local
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                                                                                              Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
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                                                                                                                                                                                                                                                 Olek A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           peptide
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                                            Claim 1; SEQ ID NO 52034; 29pp + Sequence Listing;
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This invention

describes novel oligonucleotide primers or peptide nucleic

Sequence 13 BP; 1

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8 T; 0 U; 0 Other;

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RESULT 58
ABC52018/c
ID ABC52018/c
XX ABC520
XX Oligon
XX SNP; B
XX Peptid
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                                                  This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; SEQ ID NO 52035; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          designed to detect methylation status.
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Pred. No. 3
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cardiovascular; primer;
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                                                                                                                                                                                                       This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for disgnosts and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                              Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
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ABH12189 standard; DNA; 13

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RESULT 61
ABC34878/c
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                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
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                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                      Score 10.4;
Pred. No. 3
                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                          for detecting SNP TSC0011082
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
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WPI; 2001-657177/75
                                                      Olek A,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and
                                                                                                                                                                                                                                                                                                                       This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; SEQ ID NO 116558; 29pp + Sequence Listing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       methylation
                                                                                                                                           Sequence 13
                                  2076 CCATCCAAAGAA 2087
 2 CCATCCAAAAAA 13
                                                                                       Similarity
                                                                                                                                           BP; 7 A; 4 C; 0 G; 2 T; 0 U; 0 Other;
                                                                      Conservative
                                                                                         20.8%;
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                                                                                       Score 10.4;
Pred. No. 39;
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ABF75305
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AC ABF75
XX 22-FE
XX SNP;
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XW Depti
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XX Clain
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CC acidd
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CC range SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; peptide nucleic acid; cytosine methylation; cardiovascular; primer; central nervous system; gastrointestinal; respiratory; immune; metal WO200177384-A2. Oligonucleotide 22-FEB-2002 ABF75305; ABF75305 standard; DNA; 13 18-OCT-2001 (first ij entry) NO 175302 for detecting ₽P SNP TSC0043563 metabolic. CNS;

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS ₽Ġ.

Piepenbrock C, Berlin ~

WPI; 2001-657177/75

Set of oligonucleotides, useful for designed to detect single-nucleotide methylation status. useful for diagnosis and cell e-nucleotide polymorphisms and l typing, : d cytosine

Claim 1; SEQ ID NO 175302; 29pp + Sequence Listing;

German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The

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RESULT 64
ABF30114/c
ID ABF30114 standard; DNA; 13
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                                                                                                      range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                             This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                         Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
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                                                           Sequence 13
                                                                                                                                                                                                                                                                                           Claim 1; SEQ ID NO 130111; 29pp + Sequence Listing; German.
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                                                                                          ftp.wipo.int/pub/published_pct_sequences
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Pred. No. 39;
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RESULT 66
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ID ABF443
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AC ABF443
AC ABF443
XX
DT 21-FEB
XX.

standard; DNA; 13

BP.

21-FEB-2002 ABF44362; ABF44362

(first entry)

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RESULT 65
ABF3011.5
ID ABF30
XX ABF3011.5
AC ABF30
XX ABF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                Sequence 13 BP; 9 A; 3 C; 0 G; 1 T; 0 U; 0 Other;
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AACAAACACTAA 12
                                                                                                                        Conservative
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss;
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 06-APR-2001; 2001WO-IB000713.
                                                                                                                                                                                               Oligonucleotide
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Pred. No. 39
                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                        1; cardiovascular; primer; ss;
respiratory; immune; metabolic.
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Matches 11
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                         Set of oligonuclectides, useful for diagnosis and cell typing, i designed to detect single-nuclectide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; peptide nucleic acid; cytosine methylation; central nervous system; gastrointestinal; re
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Claim 1; SEQ ID NO 52036; 29pp + Sequence Listing; German.

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Best Local Similarity
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99389, ABF00010-ABF99389, ABF000
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91.7%;
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Matches 11
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Best Local S
Matches 11
                                                                                      Sequence 13
                                                                Sequence
                                                                                                                                                                                        Claim 1; SEQ ID NO 61963; 29pp + Sequence Listing; German
                                                                                                                                                                                                         designed to detect methylation status.
                                                                                                                                                                                                                Set of oligonucleotides, useful for odesigned to detect single-nucleotide
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                                                                                                                                                                                                                                                                                                           06-APR-2001; 2001WO-IB000713
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                                                                                                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss;
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                                        Score 10.4; DB 1; Length 13; Pred. No. 39;
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ID ABF26456 standard; DNA; 13
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                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
SNP; single nucleotide polymorphism; human; diagnosis; PNA; peptide nucleic acid; cytosine methylation; cardiovascular; central nervous system; gastrointestinal; respiratory; immur
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                           Oligonucleotide SEQ ID NO 144360 for detecting SNP TSC0036296.
                                                                    21-FEB-2002
                                                                                          ABF44363;
                                                                                                                ABF44363 standard; DNA; 13 BP
                                                                                                                                                                                                                                                             Sequence 13 BP; 2 A; 0 C; 3 G; 8 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                was obtained in electronic format from W:
ftp.wipo.int/pub/published_pct_sequences
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Best Local S
Matches 11
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                                                                                                                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                          07-APR-2000; 2000DE-01019173.
                                                                                                                         06-APR-2001; 2001WO-IB000713.
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Pred. No. 39;
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RESULT 74
ABC52016/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 -ABC99989, ABF00010-ABF99989, ABH00010-ABF99989, ABH00010-ABF99989, ABH00010-ABF99989, ABH00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF999989, ABF00010-ABF999989, ABF000100-ABF999989, ABF00010-ABF999989, ABF00010-ABF999989, ABF00010-ABF999999
Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
                                                                                                                                                                                            Olek A,
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ide nucleic acid; cytosine methylation; cardiovascular;
                                                                                                                               2001-657177/75
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                                                                                                                                                                                            Piepenbrock C,
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The

Query Match

13

BP; 8 A;

3 C; 0 20.8%;

G; 2 T; 0 U;

0 Other;

Score 10.4;

DB 1; Length 13;

Claim 1;

ID NO 52033;

29pp + Sequence Listing;

German

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RESULT 75
ABH42205
ID 42045
AC ABH42
XX ABH42
XX ABH42
XX Oligo
XX SNP;
KW Pepti
KW Centr
XX Peptono
PN WO200
XX Homo
XX WO200
XX Of-AF
XX WPI;
XX WPI;
XX WPI;
XX WPI;
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                                                                                                                                                                                                                                                                                                           This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
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                                                                                     range of diseases including immune system, gastrointestinal, respirator central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC000 -ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, buwas obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1; SEQ ID NO 242182;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       designed to methylation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine
                                                       was obtained in electronic format from W
ftp.wipo.int/pub/published_pct_sequences
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Pred. No. 39;
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                                                                                                                                            Query Match
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Matches 11; Conserv
                                                                                                                                                                                                                    This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NoTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                         Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
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ABH41536;
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                                                                                          AAAAAACACTAA 2
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91.7%;
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RESULT 78
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                                                                                                                                                                                                                                                                   was obtained in electronic format from W
ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; SEQ ID NO 241513; 29pp + Sequence Listing; German.
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Pred. No. 3
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SNP; single nucleotide polymorphism; human; peptide nucleic acid; cytosine methylation; central nervous system; gastrointestinal; re

diagnosis; PNA; cancer; CNS; cardiovascular; primer; ss;

metabolic

SNP TSC0062521.

WO200177384-A2 Homo sapiens Oligonucleotide SEQ ID NO 256767 for detecting

22-FEB-2002

(first entry)

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Best Local S
Matches 11
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                                                                                                                                                                                                                                                                                                               Oligonucleotide SEQ ID NO 126454 for detecting SNP TSC0031640
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
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Set of oligonucleotides,
                               WPI; 2001-657177/75
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range of diseases including immune system, gastrointestinal, responderal nervous system, cardiovascular and metabolic disorders. I oligomers are also used for detecting cell type differentiation.

respiratory, ABC00010

ABF00010-ABF99989,

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a

Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.

SEQ ID NO 95326; 29pp + Sequence Listing; German.

WPI; 2001-657177/75.

Olek A,

Piepenbrock C,

Berlin

7.

(EPIG-)

EPIGENOMICS

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07-APR-2000; 2000DE-01019173. 06-APR-2001; 2001WO-IB000713.

Claim 1;

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RESULT 80
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ID ABC95309
AC ABC55
XX ABC95
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Pred. No. 3:
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                                                                                                                                                                                                       represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, by was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Olek A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Oligonucleotide SEQ ID NO 125414 for detecting SNP TSC0031349.
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                                                                                                                                                  Sequence 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; SEQ ID NO 125414; 29pp + Sequence Listing; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  methylation status.
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primer;
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RESULT 82
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ID ABC90
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XX SNP,
XW SNP,
XW Pepti
XW Centi
XX Centi
XX Homo
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RESULT 83
ABC95308/c
ID ABC953
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation, ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                          Oligonucleotide SEQ ID NO 95325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                   21-FEB-2002
                                                                                                                                            ABC95308
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; SEQ ID NO 90406; 29pp + Sequence Listing; German
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RESULT 84
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ID ABC05733
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
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                                                                                          Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
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Claim 1; SEQ ID NO 175301; 29pp + Sequence Listing; German

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                                                                           This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABC9989, ABF00010-ABC99899, ABF00010-ABC99899, ABF00010-ABC99899, ABF00010-ABC9989, ABF00010-ABC99899, ABF00010-ABC99899, ABF00010-ABC99899, ABF00010-ABC99899, ABF00010-ABC99899, ABF00010-ABC998999, ABF00010-ABC99899, ABF00010-ABC99989, ABF00010-ABC99989, ABF00010-ABC99989, ABF00010-ABC99989, ABF00010-ABC99989, ABF00010-ABC99989, ABF00010-ABC99989, ABF00010-ABC99989, ABF00010-ABC999
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                                                                                                                                                 Length 13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
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                                                                                                     Gaps
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RESULT 88 ABC05732/c

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RESULT 89
ABC61947
ID ABC61
XX
AC ABC61
XX
DT 21-FE
XX
Olige
XX
SNP;
KW SNP;
KW pept:
XX
OS Homo
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Best Local (
                                                                                                                                                                                                                                                                                                                                Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                    This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for disgnosts and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989, and MDIGOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Olek A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                                                                                                                                                                                                                                                                                                                             Sequence 13 BP; 2 A; 0 C; 2 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; SEQ ID NO 5723; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-657177/75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-APR-2000; 2000DE-01019173.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oligonucleotide
Homo sapiens
                                                                                            Oligonucleotide
                                                                                                                         21-FEB-2002
                                                                                                                                                                                      ABC61947 standard; DNA; 13 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABC05732 standard; DNA; 13 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (EPIG-)
                                                                                                                                                                                                                                                                                                2083 AAGAAACACTAA 2094
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                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Piepenbrock C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                        (first entry)
                                                                                            SEQ ID NO 61964 for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO
                                                                                                                                                                                                                                                                                                                                              20.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Berlin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5723 for detecting SNP TSC0001868
                                                                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                                                                                 Score 10.4;
Pred. No. 3:
                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                        detecting
                                                                                                                                                                                                                                                                                                                                                 39;
                                                                                                                                                                                                                                                                                                                                                                DB 1;
                                                                                            SNP TSC0016466.
                                                                                                                                                                                                                                                                                                                                1,
                                                                                                                                                                                                                                                                                                                                                              Length 13;
                                                                                                                                                                                                                                                                                                                                Indels
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ABH12188/c
ID ABH121
XX
AC ABH121
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF99999, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI;
                                                                                                                                           07-APR-2000; 2000DE-01019173
                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                               central nervous
                                                                                                                                                                                                                                                                                                                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Oligonucleotide SEQ ID NO 212165 for detecting SNP TSC0001691
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABH12188;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABH12188 standard; DNA; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; SEQ ID NO 61964; 29pp + Sequence Listing; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                designed to methylation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Olek A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-APR-2000; 2000DE-01019173
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11; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AACACTAAACTC 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20.8%;
ilarity 91.7%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                               system; gastrointestinal;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; 4 C; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Berlin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 10.4; D
Pred. No. 39;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                  respiratory;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    immune; metabolic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cytosine
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Olek A,

Piepenbrock C,

Berlin K;

(EPIG-)

EPIGENOMICS

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Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine methylation status.

This invention describes novel oligonucleotide primers or peptide nucleic

Claim 1; SEQ ID NO 212165; 29pp + Sequence Listing; German

acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for disgnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published\_pct\_sequences

Sequence

13

BP; 1 A; 0 C; 4 G;

8 T;

0 U;

0 Other; DB 1;

Similarity

20.8%;

Score 10.4; Pred. No. 39;

Length

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RESULT 91
ABF03882/c
ID ABF0388
XX ABF038
XX ABF038
XX Oligon
XX SNP; s
KW Peptid
KW Centra
XX Homo s
XX WO2001
XX Of-APR
XX O
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Best Local S
Matches 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Olek A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; peptide nucleic acid; cytosine methylation; cardiovascular;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oligonucleotide
                          This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                           Claim 1; SEQ ID NO 103879; 29pp + Sequence Listing;
                                                                                                                                                                                                                                                       methylation status.
                                                                                                                                                                                                                                                                                  Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-APR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        central
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2000DE-01019173
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including
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IJ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Berlin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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   system,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                              German.
                                                                                                                                                                                                                                                                                                                          typing,
                                                                                                                                                                                                                                                                                        cytosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
      respiratory
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Query Match Best Local : Matches

Similarity

20.8%;

Score 10.4; Pred. No. 3

В

1;

Length 13;

0

Gaps

0

Conservative

0

Mismatches

Sequence 13

BP; 8

A; 3 C; 0 G; 2 T; 0 U; 0 Other;

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RESULT 92
ABF61405
ID ABF6
XX ABF6
XX ABF6
XX ABF6
XX ABF6
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Matches 11
                                              This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cycosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for disgnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABCC0010-ABC9989, ABP00010-ABP9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; peptide nucleic acid; cytosine methylation; cardiovascular; primer; central nervous system; gastrointestinal; respiratory; immune; metak
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            designed to detect methylation status
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID
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Pred. No. 3
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RESULT 93
ABC90388/c
ID ABC903
 RESULT 94
ABF16560/c
ID ABF165
XX
AC ABF165
XX
DT 21-FEB
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                                                                                                                                                                                                                                                           Sequence 13 BP; 4 A; 0 C; 4
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21-FEB-2002
                              ABF16560;
                                                           ABF16560 standard; DNA; 13
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(first entry)
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                                                                                                                                                                                                                 20.8%;
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                                                                                                                                                                                                                                                              G; 5 T; 0 U; 0 Other;
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Pred. No. 3
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RESULT 95
ABF25416/c
ID ABF254
XX ABF254
XX ABF254
XX ABF254
XX Oligon
XX SNP; B
KW SNP; B
KW peptid
KW centra
XX Homo B
XX Homo B
XX Homo B
XX Homo B
XX P
PN WO2001
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                                                                                  WO200177384-A2
                                                                                                                                                                                    central nervous
                                                                                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss;
                                                                                                                                                                                                                                                                                          Oligonucleotide SEQ ID NO 125413 for detecting SNP TSC0031349
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                                                                                                                                Homo sapiens.
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                                                                                                                                                                                  system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 10.4;
Pred. No. 39;
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RESULT 96
ABH41537
ID ABH41
XX SNP;
XX SNP;
XX SNP;
XX Cent
XX Cent
XX Cent
XX Hom
XX WO2
XX Hom
XX WO2
XX Hom
XX PP 06.
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XX PP 06.
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XX O2
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Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, by was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for disgnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABCC0010 -ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073
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21-FEB-1994;
23-DEC-1994;
                            Attenuation of pathogenic HIV-1 strain NL4-3 involves deletion of 1 or more decanucleotides (AAQ96406-Q97018) from the nef gene and/or 1 or more decanucleotides (AAQ97019-Q97166) from the LTR region; the sequence of AAQ96406 corresponds to nucleotides 1-10 of the nef gene (AAQ9641). The resulting avirulent HIV strains are still capable of inducing an immune response in humans, and enable the generation of therapeutic, diagnostic and targeting agents against HIV-1 infection. (Updated on 16-OCT-2003 to
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                                                                                                                                                                                                                                                                                                                                 New non-pathogenic HIV-1 strain carrying a deletion in : LTR region - can be used in a vaccine to inhibit/reduce infection in an individual by a pathogenic strain.
   and targeting agents against standardise OS field)
                                                                                                                                                                                                                                                                         Claim 13; Page 193; 301pp; English.
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21-FEB-1994;
23-DEC-1994;
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                                                                                                                                                                                                  2094 AGCTCTCTGG 2103
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(first entry)
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94AU-00004002.
94AU-00000284.
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AAZ82752 standard; DNA; 10 BP

RESULT 100

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                                                                                                                                                     CC that are preferentially transcribed in the metastatic breast tumour cells). AAZ83941 represent tags corresponding to distinct transcripts tissue (i.e. are upregulated in metastatic breast tumour cells). AAZ83942 CC to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour cells). These CC transcripts can be used for diagnosis, prognosis, monitoring and CC treatment of breast cancer, particularly where metastatic Diagnosis is CC tompounds that modulate expression of the transcripts are potentially CC useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in selected cell types, of c.g. therapeutic genes (also ribozymes or antisense sequences), CC particularly an antigen-encoding sequence for use in gene or cell-based captures, for diagnosing breast cancer and for raising specific captus. Host cells that produce the polypeptides can be used to expand cc and isolate populations of educated, antigen-specific immune effecter cc immunotherapy.
                                                     Query Match
Best Local S
Matches 10
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19-JUN-1998;
19-JUN-1998;
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19-JUN-1998;
                                                                                                                 Sequence 10 BP; 2 A; 3 C; 1 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                treatment of cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Isolated polynucleotides differentially expressed between metastatic non-metastatic breast cancer cells, useful for diagnosis, prevention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2000-106079/09.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Metastatic breast tumour cell upregulated transcript tag #1986
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                          2092 TAAGCTCTCT 2101
                                                       1 Similarity
10; Conserv
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98US-0089997P.
98US-0090039P.
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RESULT 101
ABV65107/c
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(Mi) Is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (EST) of the invention
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immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatit
psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
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                    Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                                                                      Human skin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 10;
; Pred. No.
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34;
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Petersohn D,

Conradt M,

Hofmann

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WPI;

2002-590638/63

In vitro identification of skin-expressed

genes, useful for determining

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RESULT 103
ADK13992
ID ADK139
XX ADK139
XX ADK139
XX ADK139
XX Human
DE Human
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Best Local Similarity
Matches 10; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin The roseant is contacted.
                                                                                                                                                                                         Screening human for Rett syndrome comprises detecting mutation acid sequence encoding methyl-CpG-binding protein 2 (MECP2).
comprising detecting a mutation in a nucleic acid sequence methyl-Cpg-binding protein 2 (MECP2). The method is useful a human for Rett syndrome. The method is useful for screen:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-SEP-2000; 2000US-00657013
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        human; Rett syndrome; methyl-CpG-binding protein 2; MECP2; neurodevelopmental disease; autiem; non-syndromic mental ridiopathic neonatal encephalopathy; idiopathic infantile sidiopathic cerebral palsy; Angelman syndrome; schizophreni
                                                                                  The invention relates to a method of screening a human for Rett syndrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-MAR-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human methyl-CpG-binding protein
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                                                                                                                                                                                                                                                                                                                                                                                                 (BAYU ) BAYLOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     homeostasis and ide.g. skin cancer.
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                                                                                                                                                                                                                                                                                                                                            Veyver IB,
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                                                                                                                                             94;
                                                                                                                                          125pp; English
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  non-syndromic mental retardation; idiopathic infantile spasm;
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34;
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                                                                                                                                                                                                                                                                                                                                            Francke
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for screening
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RESULT 104
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ID ADQ357
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Best Local
                                                                                                                                                                                                                             This invention describes a novel in vitro method for identifying genes that are significant for hair-bearing skin, a first mixture of comprises recovering, from hair-bearing skin, a first mixture of genetically expressed (transcribed and optionally translated) factors (i.e. proteins, mRNA or their fragments), recovering a second, similar mixture from skin on which hair does not grow and subjecting both mixtures to serial analysis of gene expression (SAGE) to identify those genes for which expression is markedly different between the two types oskin. The invention also describes in vitro methods for determining homeostasis of human hair-bearing skin and for determining activity of cosmetic and pharmaceutical agents for use against disorders or disturbances of the homeostasis of human hair-bearing skin. A blochip and disturbances of the homeostasis of human hair-bearing skin. A blochip and
                           a test kit comprising a solid support (flexible or rigid) with immobilised probes are also described for determining homeostasis. The hair-bearing skin is from the scalp and the other skin is from the face. The method allows identification of as many as possible of the genes important for hair-bearing skin, and therefore, of a very wide range of potential therapeutic and cosmetic agents. ADQ35184-ADQ35518 represent human DNA Tag fragments used to identify genes associated with hair-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               In vitro identification of genes important for hair-bearing skin, for assessing homeostasis and in screening for pharmaceutical or agents, based on differential expression analysis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mental retardation, idiopathic neonatal encephalopathy, idiopat
infantile spasms, idiopathic cerebral palsy, Angelman syndrome
schizophrenia. The present sequence represents a mutation in the
methyl-CpG-binding protein 2, MECP2, DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 5;
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Conradt M, F
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homeostasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human hair-bearing skin-associated DNA fragment SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-SEP-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 561; 250pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AGAGCAAAAG 2068
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cosmetic; pharmaceutical;
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Pred. No.
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34;
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                                                     Matches 10;
                                                                                                                                This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 11
                                                                                                                                                                                                                                                                                                                                                                 Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss;
                                                                                                                                                                                                                                                                                                                        Claim 1; SEQ ID NO 367907; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-APR-2000; 2000DE-01019173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-APR-2001; 2001WO-IB000713.
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                         2085 GAAACACTAA 2094
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                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                acia; cycosine methylation; cardiovascular; primer; ss; system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            primer SEQ ID NO 367907 for detecting SNP TSC0056643.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Α;
                                                                                                          1 C; 2 G; 6 T; 0 U; 0 Other;
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                                                                  DB 1;
39;
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                                                                              Length 12;
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RESULT 106

Search completed: November

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2004, 15:24:46

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ADF78682/c
ID ADF786
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AC ADF786
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                                                                                                                                                                                                                                                                                                      This invention relates to a novel method of detecting chromosomal capanormalities by determining the sequence of alleles of a locus of comparing to amounts of alleles at a different, selected heterozygous locus (for example on another chromosome or a maternal locus); relative camounts are expressed as a ratio indicating presence or absence of the camounts are expressed as a ratio indicating presence or absence of the camounts are expressed as a ratio indicating presence or absence of the camounts are expressed as a ratio indicating presence or absence of the camounts are expressed as a ratio indicating presence or absence of the camounts are expressed as a ratio indicating presence or absence of the camounts are expressed as a ratio indicating presence or absence of the camounts for example translocations characters, respectively. The method is useful for transversions, monosomies, comparing the sequence and copy of comparing the sequence, and rearrangements. The camous abnormalities simultaneously, for example if several SNPs are associated with a particular disease. The method provides a rapid, non-comparing from a pregnant female, for example to detect genetic disorders as above or to determine if a foetus is a carrier of a disease or predisposed to a disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Detection of chromosomal abnormalities e.g. Down's Syndrome, non-invasively in a fetus, comprises forming a ratio of amounts of alleles a locus of interest and a different heterozygous locus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-MAR-2002; 2002US-0360232P.
11-MAR-2002; 2002US-00093618.
08-MAY-2002; 2002US-0378354P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          chromosomal abnormality; maternal locus; genetic mutation; translocation; transversion; monosomy;
                                                                                                                                                                                                                                                       Sequence 12 BP; 4 A; 3 C; 1 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 11; Page 241; 164pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-845073/78.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (DHAL/) DHALLAN R.
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Job time : 1 secs

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GENERAL INFORMATION:
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Copyright GenCore version 5.1.6 (c) 1993 - 2004 Compugen Ltd.

OM nucleic nucleic search, using 8 model

Run on: November 8, 2004, 15:27:48; ; Search time 0.001 Seconds (without alignments)
37.800 Million cell updates/s cell updates/sec

Title: Perfect score: us-10-655-847-18 50

Sequence:

Scoring table:

1 ttcagagcaaaagacttgag.....aaacactaagctctctgggc 50

IDENTITY\_NUC Gapop 10.0 , Gapext 0.5

Searched: 23 seqs, 378 residues

Total number of hits satisfying chosen parameters:

46

Minimum Maximum DB DB seq g length: 8 length: 80

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 23 summaries

Database : rnpbdb:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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Total or Company	<u>ن</u>	17,	323,	12,	189,		494,	196,	195,		381		218	217	73,	72,	71,	219,	•		73,	72,	Sequence 71, Appl	Description

RESULT 3 US-10-160-807-73/c

Sequence 73, Application US/10160807
Publication No. US20030224514A1
GENERAL INFORMATION:
APPLICANT: William Gaarde
APPLICANT: Susan M. Freier
APPLICANT: Susan M. Freier
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF PPAR-DELTA EXPRESSION
FILE REFERENCE: RTS-0189
CURRENT APPLICATION NUMBER: US/10/160,807
CURRENT APPLICATION UMBER: US/10/160,807
CURRENT FILING DATE: 2002-05-31
NUMBER OF SEQ ID NOS: 296
SEQ ID NO 73
SEQ ID NO 73

ORGANISM: Artificial Sequence

TYPE: DNA

LENGTH: 20

FEATURE:

## ALIGNMENTS

RESULT 1 US-10-160-807-71/c Sequence 71, Application US/10160807 Publication No. US20030224514A1

> Sequence 72, Application US/10160807
>
> Publication No. US20030224514A1
>
> GENERAL INFORMATION:
> APPLICANT: William Gaarde
> APPLICANT: Susan M. Freier
> APPLICANT: Susan M. Freier
> APPLICANT: Andrew T. Watt
> TITLE OF INVENTION: ANTISENSE MODULATION OF PP.
> FILE REFERENCE: RTS-0189
>
> CURRENT APPLICATION NUMBER: US/10/160,807
>
> CURRENT FILING DATE: 2002-05-31
>
> NUMBER OF SEQ ID NOS: 296
>
> SEQ ID NO 72
>
> LENGTH: 20 В Ś 밁 ફ US-10-160-807-72 US-10-160-807-71 US-10-160-807-72/c RESULT 2 FILE REFERENCE: RTS-0189
> CURRENT APPLICATION NUMBER: US/10/160,807
> CURRENT FILING DATE: 2002-05-31
> NUMBER OF SEQ ID NOS: 296
> SEQ ID NO 71
> LENGTH: 20 Matches Query Match Best Local S Matches Query Match Best Local Similarity APPLICANT: William Gaarde APPLICANT: Susan M. Freier APPLICANT: Andrew T. Watt TITLE OF INVENTION: ANTISENSE MODULATION OF PPAR-DELTA EXPRESSION TYPE: DNA ORGANISM: Artificial Sequence FEATURE: ORGANISM: Artificial Sequence FEATURE: OTHER INFORMATION: Antisense Oligonucleotide TYPE: DNA OTHER INFORMATION: Antisense Oligonucleotide 2073 GAGCCATCCAAAGAAACACT 2092 2056 TTCAGAGCAAAAGACTTGAG 2075 20 GAGCCATCCAAAGAAACACT 20 TTCAGAGCAAAAGACTTGAG 20; Similarity Conservative Conservative 40.0%; Score 20; 100.0%; Pred. No. 40.0%; Score 20; 100.0%; Pred. No. 0, 0 Mismatches Mismatches DB 1; 0.14; DB 1; Length 20; 0.14; PPAR-DELTA EXPRESSION 0 0, Length 20; Indels Indels 0; 0 Gaps Gaps 0 0

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RESULT 6
US-10-160-807-219
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US-10-160-807-217
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: Antisense Oligonucleotide US-10-160-807-73
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Best Local Similarity
Matches 20; Conserv
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CURRENT APPLICATION NUMBER: US/10/160,807
CURRENT FILING DATE: 2002-05-31
NUMBER OF SEQ ID NOS: 296
SEQ ID NO 218
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Publication No. US20030224514A1
GENERAL INFORMATION:
APPLICANT: William Gaarde
APPLICANT: Susan M. Freier
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF PPAR-DELTA EXPRESSION
FILE REFERENCE: RTS-0189
                                                                                                                     Query Match
Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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LENGTH: 20
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APPLICANT: Susan M. Freier
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF PPAR-DELTA EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/160,807
CURRENT FILING DATE: 2002-05-31
NUMBER OF SEQ ID NOS: 296
                                                                                                                                                                                                    TYPE: DNA
ORGANISM: H. sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: H. sapiens
                                                                                                                                                                                                 FEATURE:
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                                                                                          2073 GAGCCATCCAAAGAAACACT 2092
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                                                                                                                                    Score 20;
Pred. No.
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                                                                                                                        Mismatches
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                                                                                                                                      DB 1;
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0.14;
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; Sequence 219,

Application US/10160807

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Sequence 71. Application US/10655847

Publication No. US20040063129A1

GENERAL INFORMATION:
APPLICANT: William Gaarde
APPLICANT: Susan M. Freier
APPLICANT: Susan M. Freier
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF PPAR-DELTA EXPRESSION
FILE REFERENCE: RTS-0189
CURRENT APPLICATION NUMBER: US/10/655,847
CURRENT FILING DATE: 2003-09-05
PRIOR APPLICATION ONMERS: US/10/160,807
PRIOR FILING DATE: 2003-09-05
INTMBER OF SEQ ID NOS: 296
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APPLICANT: SUBAN M. Freier
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF PP.
FILE REFERENCE: RTS-0189
CURRENT APPLICATION NUMBER: US/10/655,847
CURRENT FILING DATE: 2003-09-05
PRIOR APPLICATION NUMBER: US/10/160,807
PRIOR FILING DATE: 2003-09-05
NUMBER OF SEQ ID NOS: 296
                                                                                                                                                                                                                                                         US-10-655-847-72/c
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                                                                                                                                                                      Sequence 72, Application US/10655847
Publication No. US20040063129A1
GENERAL INFORMATION:
APPLICANT: William Gaarde
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Best Local Similarity
Matches 20; Conserv
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Best Local Similarity
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CURRENT FILING DATE: 2002-05-31
NUMBER OF SEQ ID NOS: 296
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APPLICANT: Susan M. Freier
APPLICANT: Andrew T. Watt
TITLS OF INVENTION: ANTISENSE MODULATION OF PPAR-DELTA EXPRESSION
FILE REFERENCE: RTS-0189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA ORGANISM: Artificial Sequence
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100.0%; Pred. No.
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100.0%; Pred. No.
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0.14;
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0.14;
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SEQ ID NO 72 LENGTH: 20 TYPE: DNA

ORGANISM: Artificial Sequence

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RESULT 17
US-10-655-847-217
; Sequence 217, Application US/10655847
; Publication No. US20040063129A1
; GENERAL INFORMATION:
   APPLICANT: William Gaarde
; APPLICANT: SUBAN M. Freier
APPLICANT: SUBAN M. Freier
APPLICANT: SUBAN M. Freier
APPLICANT: ANDERS MODULATION OF PPAR-DELTA EXPRESSION
FILE REFERENCE: RTS-0199
CURRENT APPLICATION NUMBER: US/10/655,847
; CURRENT APPLICATION NUMBER: US/10/655,847
; PRIOR APPLICATION STATE 2003-09-05
PRIOR PILING DATE: 2003-09-05
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US-10-655-847-73/c
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; OTHER INFORMATION: Antisense Oligonucleotide
US-10-655-847-72
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APPLICANT: Susan M. Freier
APPLICANT: Susan M. Freier
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF PP.
FILE REFERENCE: RTS-0189
CURRENT APPLICATION NUMBER: US/10/655,847
CURRENT FILING DATE: 2003-09-05
PRIOR APPLICATION NUMBER: US/10/160,807
PRIOR FILING DATE: 2003-09-05
                                                                                                                            NUMBER OF SEQ ID NOS:
SEQ ID NO 217
LENGTH: 20
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SEQ ID NO 73
LENGTH: 20
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Publication No. US20040063129A1
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Query Match
Best Local Similarity
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Best Local Similarity
Matches 20; Conserv
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                                                                                   TYPE: DNA
ORGANISM: H. sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
ORGANISM: Artificial Sequence
                                                                          FEATURE:
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100.0%; Pr
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; Pred. No.
Score 20;
Pred. No.
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DB 1;
0.14;
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RESULT 11
US-10-655-847-218
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Susan M. Freier
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF PPAR-DELTA EXPRESSION
FILE REFERENCE: RTS-0189
CURRENT APPLICATION NUMBER: US/10/655,847
CURRENT FILING DATE: 2003-09-05
PRIOR APPLICATION NUMBER: US/10/160,807
PRIOR PEPLICATION NUMBER: US/10/160,807
PRIOR FILING DATE: 2003-09-05
NUMBER OF SEQ ID NOS: 296
SEQ ID NOS: 296
SEQ ID NOS: 296
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APPLICANT: William Gaarde
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SEQ ID NO 218
LENGTH: 20
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APPLICANT: William Gaarde
APPLICANT: Susan M. Freie
APPLICANT: Andrew T. Watt
                                                                                                                                              Matches
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Best Local Similarity
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APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF PPAR-DELTA EXPRESSION
FILE REFERENCE: RTS-0189
                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: H. sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: H. sapiens
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100.0%; Pred. No.
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100.0%; Pred. No.
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RESULT 13 US-10-138-674-3813 ; Sequence 3813, Application US/10138674 ; Publication No. US20040077565A1

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US-10-287-949A-3813
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US-10-138-674-3813
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                                                                                                                                                                  Sequence 195, Application US/09771933 Publication No. US20030023387A1
                                                                                                                                                 GENERAL INFORMATION:
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APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00-876-N (400/049)
CURRENT APPLICATION NUMBER: US/10/138,674
CURRENT FILING DATE: 2002-05-03
NUMBER OF SEQ ID NOS: 20822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                       -09-771-933-195/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Escobedo, Jaime TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions ReTITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor FILE REFERENCE: MeHB00-876-N (400/049) CURRENT APPLICATION NUMBER: US/10/287,949A CURRENT FILING DATE: 2003-04-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
CURRENT APPLICATION NUMBER: US/09/771,933
CURRENT FILING DATE: 2001-01-30
                                                    TITLE OF INVENTION: Computer-assisted Means TITLE OF INVENTION: Factors
                                                                                       APPLICANT: Gill-Garrison, Rosalynn D
APPLICANT: Martin, Christopher J
APPLICANT: Sanchez-Felix, Manuel V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 20822
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APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
                                    FILE REFERENCE: 620-130
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: RNA
ORGANISM: Mus musculus
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McSwiggen, Jim
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Pred. No. 1
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                                                                     for Assessing Lifestyle Risk
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; OTHER INFORMATION: Description of Artificial Sequence: Probe US-09-771-933-196
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                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Dhallan, Ravinder S.
TITLE OF INVENTION: METHODS FOR DETECTION
TITLE OF INVENTION: DISORDERS
FILE REFERENCE: 543312000420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT FILING DATE: 2001-01-30
NUMBER OF SEQ ID NOS: 205
SOFTWARE: PATENTIN VET. 2.1
SEQ ID NO 196
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SOFTWARE: PatentIn Ver. 2
SEQ ID NO 195
LENGTH: 13
                                                                                                                                                                                                                                                                                                         Sequence 494, Application US/10661165 Publication No: US20040137470A1
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Best Local Similarity
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Best Local
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APPLICANT: Martin, Christopher J
APPLICANT: Sanchez-Felix, Manuel V
TITLE OF INVENTION: Computer-assisted Means for Assessing Lifestyle Risk
TITLE OF INVENTION: Factors
FILE REFERENCE: 620-130
                 PRIOR APPLICATION NUMBER: US 60/378,354
PRIOR FILING DATE: 2002-05-08
PRIOR APPLICATION NUMBER: US 10/093,618
PRIOR FILING DATE: 2002-03-11
PRIOR APPLICATION NUMBER: US 60/360,232
PRIOR FILING DATE: 2002-03-01
                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/661,165
CURRENT FILING DATE: 2003-09-11
PRIOR APPLICATION NUMBER: PCT/US03/06198
PRIOR FILING DATE: 2003-02-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/771,933
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence FEATURE: OTHER INFORMATION: Description of Artificial Sequence: Probe
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APPLICATION NUMBER: PCT/US03/27308
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Pred. No. 8
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Pred. No. 8
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US 10/376,770

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TITLE OF INVENTION: Compositions and Methods for Wound TITLE OF INVENTION: Healing FILE REFERENCE: 000486.000116
CUBRENT APPLICATION NUMBER: US(10/314,322)
CUBRENT FILING DATE: 2002-12-09
PRIOR APPLICATION NUMBER: US 60/074,737
PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: US 60/097,937
PRIOR APPLICATION NUMBER: US 60/097,937
PRIOR FILING DATE: 1998-08-26
PRIOR APPLICATION NUMBER: US 60/102,051
PRIOR APPLICATION NUMBER: US 09/249,155
PRIOR APPLICATION NUMBER: US 09/249,155
PRIOR APPLICATION NUMBER: US 09/249,155
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US-09-249-155-189
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CURRENT FILING DATE: 1999-02-12
EARLIER APPLICATION NUMBER: 60/074,737
EARLIER FILING DATE: 1998-02-13
EARLIER APPLICATION NUMBER: 60/097,937
EARLIER FILING DATE: 1998-09-097,937
EARLIER APPLICATION NUMBER: 60/102,051
EARLIER APPLICATION NUMBER: 60/102,051
EARLIER FILING DATE: 1998-09-28
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SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 189, Application US/09249155 Publication No. US20030037345A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                Sequence 189, Application US/;
Publication No. US20030229911)
GENERAL INFORMATION:
APPLICANT: Heber-Katz, Ellen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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Best Local Similarity
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NUMBER OF SEQ ID NOS: 628
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 494
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Heber-Katz, Ellen
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TYPE: DNA
ORGANISM: Artificial Sequence
PEATURE:
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b. US20030229911A1
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Pred. No.
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; ORGANISM: Homo sapiens
US-10-450-797-323
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Best Local Similarity
Matches 10; Conserv
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                                                         SOFTWARE: PatentIn version 3.2
SEQ ID NO 323
LENGTH: 11
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ.ID:NO 189
LENGTH: 11
TYPE: DNA
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Best Local Similarity
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Publication No.
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APPLICANT: Conradt, Marcus
APPLICANT: Conradt, Marcus
APPLICANT: Hofmann, Kay
TITLE OF INVENTION: METHOD FOR DETERMINING SKIN STRESS OR SKIN AGEING IN VITRO
FILE REFERENCE: HENK-0041
CURRENT APPLICATION NUMBER: US/10/450,797
CURRENT FILING DATE: 2003-12-04
PRIOR APPLICATION NUMBER: PCT/EP01/15178
PRIOR APPLICATION NUMBER: PCT/EP01/15178
PRIOR FILING DATE: 2001-12-20
                                                                                                                                                                          FILE REFERENCE: HENK-0041
CURRENT APPLICATION NUMBER: US/10/450,797
CURRENT FILING DATE: 2003-12-04
PRIOR APPLICATION NUMBER: PCT/EP01/15178
PRIOR FILING DATE: 2001-12-20
                                                                                                                     PRIOR APPLICATION NUMBER: DE 101 00 121.5
PRIOR FILING DATE: 2001-01-03
NUMBER OF SEQ ID NOS: 1435
                                                                                                                                                                                                                                                                             APPLICANT: Conradt, Marcus
APPLICANT: Hofmann, Kay
TITLE OF INVENTION: METHOD FOR DETERMINING SKIN STRESS OR SKIN AGEING IN VITRO
                                                                                                                                                                                                                                                                                                    APPLICANT: Petersohn, Dirk
APPLICANT: Conradt, Marcu
APPLICANT: Hofmann, Kay
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ORGANISM: Homo sapiens
                                           TYPE: DNA
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No. US20040142335A1
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Pred. No. 14;
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Pred. No. 14;
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SEQ ID NO 17
LENGTH: 11
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                               ; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 32
; LENGTH: 11
; TYPE: DNA
; ORGANISM: Ehrlichia risticii
US-10-055-536-32
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Publication No. US20040191262A1
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Best Local Similarity 90.9%;
Matches 10; Conservative
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Query Match
Best Local Similarity
                                                                                                                                                                                             APPLICANT: DUTTA, Sukanta K.
APPLICANT: BISWAS, Biswajit
APPLICANT: WEMULAPALLI, Ramesh
TITLE OF INVENTION: A SIZE-VARIABLE STRAIN-SPECIFIC PROTECTIVE ANTIGEN FOR
TITLE OF INVENTION: POTOMAC HORSE FEVER
TILE REFERENCE: 8172-9016
CURRENT FPLICATION NUMBER: US/10/055,536
CURRENT FILING DATE: 2002-01-23
PRIOR APPLICATION NUMBER: US/09/157,257
PRIOR PILING DATE: 1998-09-18
PRIOR FILING DATE: 1997-09-18
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PRIOR FILING DATE: 1997-09-18
NUMBER OF SEQ ID NOS: 48
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CURRENT FILING DATE: 2002-01-23
CRIOR APPLICATION NUMBER: US/09/157,257
PRIOR FILING DATE: 1998-09-18
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APPLICANT: BISWAS, BISWAJİT
APPLICANT: VEMULAPALLI, Ramesh
TITLE OF INVENTION: A SIZE-VARIABLE STRAIN-SPECIFIC PROTECTIVE ANTIGEN FOR
TITLE OF INVENTION: POTOMAC HORSE FEVER
FILE REFERENCE: 8172-9016
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18.8%;
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Score 9.4; DB 1;
Pred. No. 14;
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Pred. No. 1
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                       Length 11;
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Matches 10; Conservative 0; Mismatches 1; Indels

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Db 1 AAAGAAATACT 11
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Search completed: November 8, 2004, 15:27:48 Job time: 0.001 secs